

# 6

## SEQUENCE LISTING



<110> Castle, Linda A.  
 Siehl, Dan  
 Giver, Lorraine  
 Minshull, Jeremy  
 Ivy, Christina  
 Chen, Yong Hong  
 Duck, Nicholas B.

<120> NOVEL GLYPHOSATE N-ACETYLTRANSFERASE  
 (GAT) GENES

<130> 02-107010US

<140> US 10/004,357  
<141> 2001-10-29

<150> US 60/244,385  
<151> 2000-10-30

<160> 515

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cattcagagc ttgaaggcga agaacagtat cagctgagag ggatggcgac gcttgaagga 240  
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aagggggcag acctttatg gtgcaatgcc aggacatctg tgagcggcta ctataaaaag 360  
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cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcggg cagtacgctt atccgccatg ccgaagagct ttttcggaaa 300  
aaggggcag acctttatg gtgcaacgcc aggacatctg tgagcggta ctataaaaaag 360  
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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca taatgccgaa 180  
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aaaggcgcgg acctttatg gtgcaaacgcc aggatatctg tgagcggcta ctatgaaaag 360  
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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca taatgccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
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cacctcggtg gatattacca gggcaagctg atcagcatcg cttcccttca taaagccgaa 180  
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taccgtgagc aaaaagcggg aagcacgctc atccgcccattg ccgaagagct tcttcggaaaa 300  
aagggggcag acctttatg gtgcaatgcc aggacatctg tgagcggcta ctatgaaaag 360  
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<211> 146  
<212> PRT  
<213> Bacillus licheniformis

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Thr Asp Leu Leu Gly Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly			
35	40	45	
Lys Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Glu Leu			
50	55	60	
Glu Gly Glu Glu Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly			
65	70	75	80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu			
85	90	95	
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr			
100	105	110	
Ser Val Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu			
115	120	125	
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys			
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Leu Thr			
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<212> PRT  
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<223> Unidentified microorganism derived from soil sample

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Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Asp
   35         40          45
Arg Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu
   50         55          60
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly
   65         70          75          80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu
   85         90          95
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr
  100        105         110
Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly
  115        120         125
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys
  130        135         140
Leu Thr
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<210> 8  
<211> 146  
<212> PRT  
<213> Unknown

<220>  
<223> Unidentified microorganism derived from soil sample

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Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Asn Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Ile  
100 105 110  
Ser Val Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Ile Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Ala  
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<210> 9  
<211> 146  
<212> PRT  
<213> Unknown

<220>  
<223> Unidentified microorgansim derived from soil sample

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Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Asn Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Ile  
100 105 110  
Ser Val Ser Gly Tyr Tyr Glu Lys Leu Gly Leu Ser Glu Gln Gly Gly  
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Ile Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Ala  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
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Thr Asp Leu Leu Gly Gly Ala Phe His Leu Gly Gly Tyr Tyr Gln Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Glu Glu Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
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cacctcggtg gatattaccg gggcaagctg atcagcatcg cctcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaagga 240  
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aaaggcgccgg accctttatg gtgcaacgccc aggacgtctg cgagcgggta ctataaaaag 360  
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cacctcggtg gatattaccg gggcaagctg gtcagcatcg cctcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aagacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcgaa cagtacgctt atccgcccattt cccgaagagct tcttcggaaa 300  
aaaggcgccgg accctttatg gtgcaacgccc aggacatctg cgagcgggta ctataaaaag 360  
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<210> 13  
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aaaggcgcgg accttttatg gtgcaacgcgc aggacatctg cgagcgggta ctataaaaag 360  
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<220>  
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aaaggcgcgg accttttatg gtgcaacgcgc aggacatctg cgagcgggta ctataaaaag 360  
ctcggttca gcaacacaggc cgaagtctac gacacaccgc cggtcggacc ccatatgg 420  
atgtataaga aattgacgtat a 441

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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagga 240  
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aaggggcag acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
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atgtataaga aattgacgta a 441

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aaggggcag acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
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aagggcgcag acctttatg gtgcaacgcc aggacatctg cgagcggta ctataaaaag 360  
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aagggcgcag acctttatg gtgcaacgcc aggacatctg cgagcggta ctataaaaag 360  
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aaggggcgcag acctttatg gtgcaacgcc aggacatctg cgagcggta ctataaaaag 360  
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catccagagc ttgaaggccg aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgcggc aaaaagcggg cagtcgtt atccgcattt 300  
aaaggcgcgg acctttatg gtgcaacgcc aggacatctg cgagcggta ctataaaaag 360  
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cacctcggtg gatattaccg gggcaagctg gtcagcatcg cttcccttca tcaagccgaa 180  
cacccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaagga 240  
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aagggggcag accttttatg gtgcaacgcg aggacatctg cgagcgggta ctataaaaag 360  
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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
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taccgcgagc aaaaagcggg cagtacgctt atccgcccattg ccgaagagact tttcgaaaa 300  
aagggggcag accttttatg gtgcaacgcg aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcaacacagg cgaagtctac gacacaccgc cggtcggacc tcataat 420  
atgtataaga aattgacgta a 441

<210> 26

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 26

atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60

cggccgaatc agccgctgga agcatgcaag tatgaaaccg atttgcttag ggggcgttt 120  
cacctcggtg gatattacccg gggcaagctg atcagcatacg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaaggg 240  
taccgtgagc aaaaagcggg aagcacgctc atccgcccattg ccgaagagct tcttcggaaa 300  
aaaggcgcgg acctttatg gtgcaacgccc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcgaaacaggg cgaagtctac gacataccgc cggtcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 27  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 27  
atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcatttcc 60  
cgccgaatc agccgctgga agcatgcaag tatgaaaccg atttgctcg gggcacgttt 120  
cacctcggtg gatattacccg gggcaagctg gtcagcatacg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcggg aagtacgctt atccgcccattg ccgaagagct tcttcggaaa 300  
aaggggccag acctttatg gtgcaacgccc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcgaaacaggg cgaagtctac gacataccgc cgaccggacc ccataatgg 420  
atgtataaga aattgacgta a 441

<210> 28  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 28  
atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcatttcc 60  
cgccgaatc agccgcttga agcatgtatg tatgaaaccg atttgctcg gggcacgttt 120  
cacctcggtg gatattacccg gggcaagctg gtcagcatacg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcggg cagtaacgctt atccgcccattg ccgaagagct tcttcggaaa 300  
aaggggccag acctttatg gtgcaacgccc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcgaaacaggg cgaagtctac gacataccgc cgaccggacc ccataatgg 420  
atgtataaga aattgacgta a 441

<210> 29  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 29  
atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcatttcc 60  
cgccgaatc agccgctgga agcatgcaag tatgaaaccg atttgctcg gggcacgttt 120  
cacctcggtg gatattacccg gggcaagctg gtcagcatacg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
taccgtgagc aaaaagcggg cagtaacgctt atccgcccattg ccgaagagct tcttcggaaa 300  
aaggggccag acctttatg gtgcaacgccc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcgaaacaggg cgaagtctac gacataccgc cggtcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 30  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 30  
atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcattctc 60  
cgccgaatc agccgcttga agcatgcaag tatgaaaccg atttgctcg gggtgcgtt 120  
cacctcggtg gatattaccc gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaagga 240  
taccgcgagc aaaaagcggg aagtacgctt atccgcctatg ccgaagagct tcttcggaaa 300  
aaaggcgcgg accctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggtttca gcgacaagg cgggtctac gatataccgc cgatcggacc tcataaaaa 420  
atgtataaga aattgacgta a 441

<210> 31  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 31  
atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcattctc 60  
cgccgaatc agccgcttga agcatgcaag tatgaaaccg atttgctcg gggtacgtt 120  
cacctcggtg gatattacca gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
taccgcgagc aaaaagcggg aagtacgctt atccgcctatg ccgaagagct tcttcggaaa 300  
aagggggcag accctttatg gtgcaacgcc aggacatctg tgagcgggta ctataaaaag 360  
ctcggtttca gcgacaagg cgggtctac gatataccgc cgatcggacc tcataaaaa 420  
atgtataaga aattgacata a 441

<210> 32  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 32  
atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcattctc 60  
cgccgaatc agccgcttga agcgtgtatg tatgaaaccg atttgctcg gggcacgtt 120  
cacctcggtg gatattacca gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
taccgcgagc aaaaagcggg cagtacgctt atccgcctatg ccgaagagct tcttcggaaa 300  
aagggggcag accctttatg gtgcaacgcc aggacatctg cgagcggcta ctatgaaaag 360  
ctcggtttca gcgacaagg cgggtctac gatataccgc cgatcggacc tcataaaaa 420  
atgtataaga aattggcata a 441

<210> 33  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 33

atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcataactc 60  
cgccgaatc agccgcttgg a gcatgcaag tatgaaaccg atttgcttag gggtcgttt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
caccaggagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcggg cagtcgctt atccgcatg ccgaagcgct tcttcggaaa 300  
aaaggcgcgg acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcaacaggg cgaagtctac gacataccgc caactggccc ccataatttg 420  
atgtataaga aattgacgta a 441

<210> 34

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 34

atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcataactc 60  
cgccgaatc agccgcttgg a gcatgcaag tatgaaaccg atttgcttag gggcacgttt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccaggagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcggg a a g c a c g c t c a t c g c a g a g g t t c t t c g g a a a 300  
a a g g g g c a g a c t t t a t g g t g c a a c g c c a g g a c a t c t g c g a g c g g g t a c t a a a a a g 360  
ctcggcttca gcaacaggg cgaagtccac gacataccgc cgaccggacc ccataatttg 420  
atgtataaga aattgacgta a 441

<210> 35

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 35

atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcatttctc 60  
cgccgaatc agccgcttgg a gcatgcaag tatgaaaccg atttgcttag gggtacgttt 120  
cacctcggtg gatattaccg gggcaagctg gtcagcatcg cttcccttca tcaagccgaa 180  
catccaggagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcggg cagtcgctt atccgcatg ccgaagaggt tcttcggaaa 300  
a a g g g g c a g a c t t t a t g g t g c a a c g c c a g g a c a t c t g c g a g c g g g t a c t a a a a g 360  
ctcggcttca gcaacaggg cgaagtctac gacacaccgc cggtcggacc tcataatttg 420  
atgtataaga aattgacgta a 441

<210> 36

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 36

atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcatttctc 60  
cgccgaatc agccgcttgg a gcatgcaag tatgaaaccg atttgcttag gggcacgttt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccaggagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaaggg 240

taccgtgagc aaaaagcggg aagtacgctc atccgccatg ccgaagagct tcttcggaaa 300  
aagggggcag acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggttca gccaacaagg cgggtctac gacataccgc cggtcggacc tcataatttg 420  
atgtataaga aattgacgta a 441

<210> 37  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 37  
atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcatttctc 60  
cgccgaatc agccgctgga agcatgcaag tatgaaacccg atttgcctgg gggcacgttt 120  
cacctcggtg gatattatcg gggcaagctg atcagcatcg cttcccccata tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagttt cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg aagcacgctc atccgcctatg ccgaagagct tcttcggaaa 300  
aagggggcag acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggttca gccaacaagg cgggtctac gacataccgc cggtcggacc tcataatttg 420  
atgtataaga aattgacgta a 441

<210> 38  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 38  
atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcatttctc 60  
cgccgaatc agccgctgga agcatgcaag tatgaaacccg atttgcctgg gggcacgttt 120  
cacctcggtg gatattatcg gggcaagctg gtcagcatcg cttcccccata tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagttt cagctgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagcggg cagtcgtt atccgcctatg ccgaagagct tcttcggaaa 300  
aagggggcag acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggttca gccaacaagg cgggtctac gacataccgc cggtcggacc tcataatttg 420  
atgtataaga aattgacgta a 441

<210> 39  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 39  
atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcatttctc 60  
cgccgaatc agccgctgga agcatgtatg tatgaaacccg atttgcctgg gggcacgttt 120  
cacctcggtg gatattatcg gggcaagctg atcagcatcg cttcccccata tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagttt cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg aagcacgctc atccgcctatg ccgaagagct tcttcggaaa 300  
aagggggcag acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggttca gccaacaagg cgggtctac gacataccgc cggtcggacc tcataatttg 420  
atgtataaga aattgacgta a 441

<210> 40  
<211> 441

<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 40  
atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgcttga agcatgtatg tatgaaaccg atttgctcg gggtacgtt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaagga 240  
taccgcgagc aaaaagcggg aagcacgctc atccgcctatg ccgaagagct tttcgaaaa 300  
aaggggcag acctttatg gtgcaacgcc aggacatctg cgagcggcta ctataaaaag 360  
ctcggcttca gccaacaggg cgaagtctac gacataccgc cgaccggacc ccataatttg 420  
atgtataaga aattgacgta a 441

<210> 41  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 41  
atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcataactc 60  
cgccgaatc agccgcttga agcatgtatg tatgaaaccg atttgctcg ggacacgtt 120  
cacctcggtg gatattaccg gggcaagctg gtcagcatcg cttcccttca ccaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcggg cagtcgcatt atccgcctatg ccgaagcgct tttcgaaaa 300  
aaaggcgcgg acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gccaacaggg cgaagtctac gacataccgc cgaccggacc ccataatttg 420  
atgtataaga aattgacata a 441

<210> 42  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 42  
atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgcttga agcatgtatg tatgaaactg atttgctcg ggacacgtt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgcgagc aaaaagcggg cagtcgcatt atccgcctatg ccgaagagct tttcgaaaa 300  
aaggggcag acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gccaacaagg cggggcttac gacataccgc cgaccggacc ccataatttg 420  
atgtataaga aattgacgta a 441

<210> 43  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 43

atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgcttga agcatgcaag tatgaaaccg atttgctcg gggcacgtt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagga 240  
taccgcgagc aaaaagcggg cagtacgctt atccgccatg ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggc cgaagtctac gacacaccgc cgatcggacc tcataat 420  
atgtataaga aattgacgta a 441

<210> 44

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 44

atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgcttga agcatgcaag tatgaaaccg atttgctcg gggtacgtt 120  
cacctcggtg gatattacca gggcaagctg atcagcatcg cttccttca taaagccgaa 180  
cattcagagc ttgagggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgcgagc aaaaagcggg cagtacgctt atccgccatg ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaatgcc aggacatctg tgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaagg cgggatctac gacataaccgc cgatcggacc tcataat 420  
atgtataaga aattgacgta a 441

<210> 45

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 45

atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgcttga agcatgcaag tatgaaaccg atttgctcg gggtacgtt 120  
cacctcggtg gatattacca gggcaagctg atcagcatcg cttccttca taaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaagga 240  
taccgtgagc aaaaagcggg aagcacactc atccgccatg ccgaagagct tcttcggaaa 300  
aaggcgcag acctttatg gtgcaacgcc aggacatctg tgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggc cgaagtctac gacataaccgc cgatcggacc tcataat 420  
atgtataaga aattgacgta a 441

<210> 46

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 46

atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgcttga agcatgcaag tatgaaaccg atttgctcg gggtgcgtt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgcgagc aaaaagcggg aagtacgctt atccgccatg ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaagg cgggatctac gacataaccgc cgatcggacc tcataat 420

atgtataaga aattgacgta a 441

<210> 47  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 47  
atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgcttga agcatgtatg tatgaaacccg atttgctcg gggtacgttt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttaa tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg aagtacgctt atccgcctg ccgaagagct tcttcggaaa 300  
aaaggcgcgg accttttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggc cgaagtctac gacacaccgc cggtcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 48  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 48  
atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgcttga agcatgtatg tatgaaacccg atttgctcg aggacacgttt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttaa tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaaggg 240  
taccgtgagc aaaaagcggg aagcacgctc atccgcctg ccgaagagct tcttcggaaa 300  
aaggggcag accttttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggc cgaagtctac gacacaccgc cggtcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 49  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 49  
atgatcgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgcttga agcatgtatg tatgaaacccg atttgctcg gagtgcggttt 120  
cacctcggcg gatattaccg gggcaagctg atcagcatcg cttcccttca ccaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgaggg ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg aagtacgctt atccgcctg ccgaagagct tcttcggaaa 300  
aaggggcag accttttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggc cgaagtctac gacacaccgc cggtcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 50  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 50

atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcattctc 60  
cgccgaatc agccgcttga a gcatgcaag tatgaaacccg atttgctcg gggcacgttt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccagac ttgaaggcca aaaacagtat cagctgagag ggtatggcgac actcgaagga 240  
taccgtggc aaaaagcggg cagtcgcctt atccgcctatg ccgaagagct tcttcggaaa 300  
aaaggcgcgg acctttgtg gtgcaacgccc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcaacacagg cgaagtctac gacataccgc cgactgggccc ccataattttg 420  
atgtataaga aattgacgta a 441

<210> 51

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 51

atgattgaag tcaaaccatat aaatgcggaa gatacgtatg agatcaggca ccgcataactc 60  
cgccgaatc agccgcttga a gcatgcaag tatgaaacccg atttgctcg gggtcgttt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccagaac ttgaaggcca aaaacagtat cagctgagag ggtatggcgac acttgaagga 240  
taccgtggc aaaaagcggg tagtacgcctt atccgcctatg ccgaagagct tcttcggaaa 300  
aaaggcgcgg acctttgtg gtgcaacgccc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcaacacaagg cgaagtctac gacataccgc cgactgggccc ccataattttg 420  
atgtataaga aattgacgta a 441

<210> 52

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 52

atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcattctc 60  
cgccgaatc agccgcttga a gcatgcaag tatgaaacccg attcgcctcg gggcacgttt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttccctttaa tcaagccgaa 180  
catccagac ttgaagggtca aaaacagtat cagctgagag ggtatggcgac acttgaagga 240  
taccgtggc aaaaagcggg cagtcgcctt atccgcctatg ccgaagagct tcttcggaaa 300  
aaaggcgcgg accttttatg gtgcaacgccc aggacgtctg cgagcgggta ctataaaaag 360  
ctcggcttca gcaacacaagg cggggcttac gacataccgc cggtcggacc tcataattttg 420  
atgtataaga aattgacgta a 441

<210> 53

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 53

atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcattctc 60  
cgccgaatc agccgcttga a gcatgcaag tatgaaacccg atctgcgtgg gggcacgttt 120  
caccttaggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180

catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg aagtacgctt atccgcccattt ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaacgcc aggacatctg cgagcggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacacaccgc cggtcggacc tcataattttg 420  
atgtataaga aattgacgta a 441

<210> 54  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 54  
atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcataactc 60  
cggccgaatc agccgctaga agcatgcaag tatgaaaccg atttgcttag gggtgcgttt 120  
cacctcggtg gatattaccg gggcaagctg atcagcattcg cttcccttca tcaagctgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg aagtacgctc atccgcccattt ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaacgcc aggacatctg cgagcggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacacaccgc cggtcggacc tcataattttg 420  
atgtataaga aattgacgta a 441

<210> 55  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 55  
atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcatttctc 60  
cggccgaatc agccgcttgg aagcatgcaag tatgaaaccg atttgcttag gggtgcgttt 120  
cacctcggtg gatattaccg gggcaagctg gtcagcattcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg aagcacgcctc atccgcccattt ccgaagagct tcttcggaaa 300  
aagggcgcag acctttatg gtgcaacgcc aggacatctg cgagcggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacacaccgc cggtcggacc tcataattttg 420  
atgtataaga aattgacgta a 441

<210> 56  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 56  
atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcataactc 60  
cggccgaatc agccgcttgg aagcatgcaag tatgaaaccg atttgcttag gggtgcgttt 120  
cacctcggtg gatattaccg gggcaagctg gtcagcattcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgtgagc aaaaagcggg cagtcgcattt atccgcccattt ccgaagagct tcttcggaaa 300  
aagggcgcgg acctttgtg gtgcaacgcc aggacatctg cgagcggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacacaccgc cggtcggacc tcataattttg 420  
atgtataaga aattgacata a 441

<210> 57

<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 57  
atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgcgaatc agccgcttga agcatgcaag tatgaaaccg atttgctcg gggcacgtt 120  
cacctcggcg gatattaccg gggcaagctg gtcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg cagtcgcattt atccgcctg ccgaagcgct tcttcggaaa 300  
aaggggcag acctttatg gtgcaacgcg aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcaacaggg cgaagtctac gacacaccgc cgaccggacc ccatattttg 420  
atgtataaga aattgacgta a 441

<210> 58  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 58  
atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgcgaatc agccgcttga agcatgtaag tatgaaaccg atttgctcg gggcacgtt 120  
cacctcggcg gatattaccg gggcaagctg gtcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagctgg cagtcgcattt atccgcctg ccgaagcgct tcttcggaaa 300  
aagggcgcgg acctttgtg gtgcaacgcg aggacatctg cgagcggcta ctataaaaag 360  
ctcggcttca gggacaagg cgggtctac gacataccgc ctgtcggacc tcatattttg 420  
atgtataaga aattgacgta a 441

<210> 59  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 59  
atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgcgaatc agccgcttga agcatgcaag tatgaaaccg atttgctcg gggcacgtt 120  
cacctcggcg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg cagtcgcattt atccgcctg ccgaagagct tcttcggaaa 300  
aagggcgcgg acctttgtg gtgcaacgcg aggacgtctg cgagcgggta ctataaaaag 360  
ctcggcttca gcaacaggg cgaagtctac gacataccgc cgactgggcc ccatattttg 420  
atgtataaga aattgacgta a 441

<210> 60  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 60  
atgattgaag tcaaaccaat aaatgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgttga agcatgcaag tatgaaaccg atttgctcg gggtacgtt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcctttaa tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaagga 240  
taccgtgagc aaaaagcggg cagtagctt atccgcatg ccgaagagct tcttcggaaa 300  
aaaggcgcgg acctttatg gtgcaacgcc aggacgtctg cgagcgggta ctataaaaag 360  
ctcggttca gcaacaagg cgggtctac gacataccgc cggtcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 61  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 61  
atgattgaag tcaaaccaat aaatgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgttga agcatgcaag tatgaaaccg atttgctcg gggtgcgtt 120  
cacctcggtg gatattaccg gggcaagctg gtcagcatcg cttccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaaggg 240  
taccgtgagc aaaaagcggg cagtagctt atccgcatg ccgaagagct tcttcggaaa 300  
aaaggcgcgg acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggttca gcaacaagg cgggtctac gacataccgc cgactggcc ccatatgg 420  
atgtataaga aattgacgta a 441

<210> 62  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 62  
atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
aggccgaatc agccgttga agcatgcaag tatgaaaccg atttgctcg gggtgcgtt 120  
cacctcggtg gatattaccg gggcaagctg gtcagcatcg cttccttca tcaggccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagcggg aagcacgctt atccgcatg ccgaagcgct tcttcggaaa 300  
aaaggcgcgg acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggttca gcaacaagg cgggtctac gacataccgc cggccggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 63  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 63  
atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgttga agcatgcaag tatgaaaccg atttgctcg gagcacgtt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaaggg 240  
taccgtgagc aaaaagcggg aagcacgctc atccgcatg ccgaagagct tcttcggaaa 300  
aaaggcgcgg acctttatg gtgcaacgcc aggacgtctg cgagcggcta ctataaaaag 360

ctcggcttca gcbaacaggg cgaagtctac gacacaccgc cggtcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 64  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 64  
atgattgaag tcaaaccat aaacgcggaa gatacgatg agatcaggca ccgcatttc 60  
cgccgcgaaatc agccgcttga agcatgtatg tatgaaaccg atttgctcg gggtacggtt 120  
cacctcggtg gatattaccg gggcaagcta gtcagcatcg ctccctttaa tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagttt cagctgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagcggtt cagtcgctt atccgcattt ccgaagagct tcttcggaaa 300  
aaaggcgccgg accctttgtt gtgcaacgc 360  
ctcggcttca gcbaacaggg cgaagtctac gacacaccgc cggtcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 65  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 65  
atgattgaag tcaaaccat aaacgcggaa gatacgatg agatcaggca ccgcatttc 60  
cgccgcgaaatc agccgcttga agcatgcaag tatgaaaccg atttgctcg gggtacggtt 120  
cacctcggtg gatattaccg gggcaagctt atcagcatcg ctccctttaa tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagttt cagctgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagcggtt cagtcgctt atccgcattt ccgaagagct tcttcggaaa 300  
aaaggcgccgg acccttttatg gtgcaacgc 360  
ctcggcttca gcbaacaagg cggggcttac gacatgccgc cggtcggacc tcataatgg 420  
atgtataaga agttgacgta a 441

<210> 66  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 66  
atgattgaag tcaaaccat aaacgcggaa gatacgatg agatcaggca ccgcatttc 60  
cgaccgcgaaatc agccgcttga agcatgcaag tatgaaaccg atttgctcg aggacacggtt 120  
cacctcggtg gatattaccg gggcaagctt atcagcatcg ctccctttaa tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagttt cagctgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagcggtt cagtcgctt atccgcattt ccgaagacgct tcttcggaaa 300  
aaggggcgac acccttttatg gtgcaacgc 360  
ctcggcttca gcbaacaggg cgaagtctac gacacaccgc cggtcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 67  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 67  
atgattgaag tcaaaccaat aaacgcggag gatacgtatg agatcaggca ccgcatttc 60  
cgcccgaaatc agccgctgga agcatgcaag tatgaaaccg atttgctcg gggtgcgtt 120  
cacctcggtg gatattaccg gggcaagctg gtcagcatcg cctccttca tcaagccgaa 180  
catccagagc ttgaaggcct aaaacagtat cagctgagag ggatggcgac actcgaaggg 240  
taccgtgagc aaaaagcggg aagtacgctc atccgcatg ccgaagagct tcttcggaaa 300  
aaggggccag acctcttatg gtgcaacgcc aggacgtctg cgagcgggta ctataaaaag 360  
ctcggtttca gcbaacaggg cgaagtctac gacacaccgc cgaccggacc tcataaaaa 420  
atgtataaga aattgacgta a 441

<210> 68  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 68  
atgattgaag tcaaaccaat aaacgcggag gatacgtatg agatcaggca ccgcatttc 60  
cgcccgaaatc agccgctgga agcatgcaag tatgaaaccg atttgctcg gggtgcgtt 120  
cacctcggtg gatattaccg gggcaagctg gtcagcatcg cctccttca tcaagccgaa 180  
catccagagc ttgaaggcct aaaacagtat cagctgagag ggatggcgac actcgaaggg 240  
taccgtgagc aaaaagcggg aagtacgctc atccgcatg ccgaagagct tcttcggaaa 300  
aaggggccag acctcttatg gtgcaacgcc aggacgtctg cgagcgggta ctataaaaag 360  
ctcggtttca gcbaacaggg cgaagtctac gacacaccgc cgaccggacc tcataaaaa 420  
atgtataaga aattgacgta a 441

<210> 69  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 69  
atgattgaag tcaaaccaat aaacgcggag gatacgtatg agatcaggca ccgcatttc 60  
cgcccgaaatc agccgctgga agcatgcaag tatgaaaccg atttgctcg gggtacgtt 120  
cacctcggtg gatattaccg gggcaagttg gtcagcatcg cctccttca tcaagccaaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaaggg 240  
taccgtgagc aaaaagcggg tagtacgctt atccgcatg ccgaagagct tcttcggaaa 300  
aaaggcgcgg accttttatg gtgcaacgcc aggacgtctg cgagcgggta ctataaaaag 360  
ctcggtttca gcbaacaggg cgaagtctac gacacaccgc cggtcggacc tcataaaaa 420  
atgtataaga aattgacgta a 441

<210> 70  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 70  
atgattgaag tcaaaccaat aaacgcagaa gatacgtatg agatcaggca ccgcatttc 60  
cgcccgaaatc agccgctgga agcatgcaag tatgaaaccg atttgctcg gggtgcgtt 120

cacctcggtg gatattaccc gggcaagctg atcagcatcg cttcccttaa tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagttgagag ggatggcgac acttgaagag 240  
taccgtgagc aaaaagcggg aagtacgctt atccgcctatg ccgaagcgct tcttcggaaa 300  
aaggggggcag acctcttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacacaccgc cggtcggacc tcataatttg 420  
atgtataaga aattgacgta a 441

<210> 71  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 71  
atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcatttctc 60  
cgccgaatc agccgcttga agcatgcaag tatgaaaccg atttgctcg gggtacgttt 120  
cacctcggtg gatattaccc gggcaagctg gtcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagcggg aagcacgctc atccgcctatg ccgaagagct tcttcggaaa 300  
aaaggcgcgg accttttgtt gtgcaacgcc aggacatctg cgagcgggta ctataaaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacacaccgc cggtcggacc tcataatttg 420  
atgtataaga aattgacgta a 441

<210> 72  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 72  
atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcatttctc 60  
cgccgaatc agccgcttga agcatgcaag tatgaaaccg atttgctcg gggtacgttt 120  
cacctcggtg gatattaccc gggcaagctg gtcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgtgagc aaaaagcggg cagtcgctt atccgcctatg ctgaagcgct tcttcggaaa 300  
aaaggcgcgg accttttgtt gtgcaacgcc aggacatctg caagcgggta ctataaaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacacaccgc cgactgggcc ccataatttg 420  
atgtataaga aattgacgta a 441

<210> 73  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 73  
atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcatttctc 60  
cgccgaatc agccgcttga agcatgcaag tatgaaaccg atttgctcg gggtgcgttt 120  
cacctcggtg gatattaccc gggcaagctg gtcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagcggg aagtacgctc atccgcctatg ccgaagagct tcttcggaaa 300  
aaggggggcag acctcttatg gtgcaacgcc aggacaactg cgagcgggta ctataaaaaag 360  
ctcggcttca gcbaacaggg tgaagtctt gacacaccgc cgaccggacc ccataatttg 420  
atgtataaga aattgacgta a 441

<210> 74  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 74  
atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgcgaatc agccgcttga agcatgtatg tatgaaaccg atttgctcg gggcacgtt 120  
caccttaggtg gatattaccg gggcaagctg gtcagcatcg cctcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg aagcacgctc atccgcccatttccgaa 300  
aagggggcag accttttatg gtgcaacgccc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cggtcggacc tcataat 420  
atgtataaga aattgacgta a 441

<210> 75  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 75  
atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgcgaatc agccgcttggaa agcatgcaag tatgaaaccg atttgctcg gggtgcgtt 120  
caccttgggtg gatattaccg gggcaagctg gtcagcatcg cctccctttaa tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaagga 240  
taccgtgagc aaaaagcggg cagtaacgctt atccgcccatttccgaa 300  
aagggggcag accttttatg gtgcaacgccc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgaccggacc ccataat 420  
atgtataaga aattgacgta a 441

<210> 76  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 76  
atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcataactc 60  
cgccgcgaatc agccgcttggaa agcatgcaag tatgaaaccg atttgctcg gggtgcgtt 120  
caccttgggtg gatattaccg gggcaagctg atcagcatcg cctcccttca tcaagccgaa 180  
cacccttgggtg gatattaccg gggcaagctg atcagcatcg cctcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcggg cagtaacgctt atccgcccatttccgaa 300  
aaggggcggg accttttatg gtgcaacgccc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc caactgggcc ccataat 420  
atgtataaga aattgacgta a 441

<210> 77  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 77  
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cgccgcgaatc agccgctgga agcatgcaag tatgaaaccg atttgctcg gggtacggtt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cctcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagcggg cagtacgctt atccgccatg ccgaagcgct tcttcggaaa 300  
aaggggcag acctttagt gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgaccggacc ccataatttg 420  
atgtataaga aattgacgta a 441

<210> 78  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 78  
atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcatttctc 60  
cgccgcgaatc agccgctgga agcatgtatg tatgaaaccg atttgctcg gggtgcgtt 120  
cacctcggtg gatattaccg gggcaagctg gtcagcatcg cctcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcggg cagtacgctt atccgccatg ccgaagagct tcttcggaaa 300  
aagggcgcgg acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacacaccgc cggtcggacc tcataatttg 420  
atgtataaga agttgacgta a 441

<210> 79  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 79  
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cacctcggtg gatattaccg gggcaagctg gtcagcatcg cctcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat caactgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagcggg cagtacgctt atccgccatg ccgaagagct tcttcggaaa 300  
aagggcgcgg acctttatg gtgcaacgcc aggacgtctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg caaagtctac gacataccgc cggtcggacc tcataatttg 420  
atgtataaga aattgacgta a 441

<210> 80  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 80  
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cgccgcgaatc agccgctgga agcatgcaag tatgaaaccg atttgctcg gggtacggtt 120  
cacctcggtg gatattaccg gggcaagctg gtcagcatcg cctcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaagag 240  
taccgcgagc aaaaagcggg aagcacgctc atccgccatg ccgaagagct tcttcggaaa 300

aagggggcag acctttatg gtgcaacgcc aggacatctg cgagcggta ctataaaaag 360  
ctcggttca gccaacaggg cgaagtctac gacataccgc cggtcggacc tcataatttg 420  
atgtataaga aattgacata a 441

<210> 81  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 81  
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caccttggcgatattaccggggcaagctgatcagcatgtttcccttca tcaagccgaa 180  
catccagagcttgaaggccaaaacagtatcagctgagggatggcgacttgaaggg 240  
taccgtgagcaaaaagcgggcagcacgctt atccgccttgcgaagagcttccggaaa 300  
aaaggcgcggacctttatgtgtcaacgcaggacatctgcgagcggta ctataaaaag 360  
ctcggttca gccaacaggg cgaagtctac gacataccgc cggtcggacc tcataatttg 420  
atgtataacgaaattgacgtta 441

<210> 82  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 82  
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cgccgaatc agccgcttga agcatgcaag tatgaaaccg atttgcgtt gggtacgtt 120  
caccttggcgatattaccggggcaagctgtcagcatgcctcccttca tcaagccgaa 180  
catccagagcttgaaggccaaaacagtatcagctgagggatggcgacttgaaggg 240  
taccgtgagcaaaaagcgggcagcacgctt atccgccttgcgaagagcttccggaaa 300  
aaaggcgcggacctttatgtgtcaacgcaggacatctgcgagcggta ctataaaaag 360  
ctcggttca gccaacaggg cgaagtctac gacataccgc cggtcggacc tcataatttg 420  
atgtataacgaaattgacgtta 441

<210> 83  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 83  
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caccttggcgatattaccggggcaagctgtcagcatgcctcccttca tcaagccgaa 180  
catccagagcttgaaggccaaaacagtatcagctgagggatggcgacttgaaggg 240  
taccgtgagcaaaaagcgggcagcacgctt atccgccttgcgaagagcttccggaaag 300  
aaaggcgcggacctttatgtgtcaacgcaggacatctgcgagcggta ctataaaaag 360  
ctcggttca gccaacaggg cgaagtctac gacataccgc cgaccggacc ccataatttg 420  
atgtataacgaaattgacgtta 441

<210> 84  
<211> 441  
<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 84

atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcattctc 60  
cgccgcatac agccgcttga agcatgcaag tatgaaaccg atttgctcg gggtacgttt 120  
cacctcggtg gatattaccg gggcaagctg gtcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg aagtacgctt atccgcctatg ccgaagagct tcttcggaaa 300  
aaaggcgcgg accttttatg gtgcaacgcg aggacatctg cgagcgggtta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgactgggcc ccataattttg 420  
atgtataaga aattgacgta a 441

<210> 85

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 85

atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcataactc 60  
cgccgcatac agccgcttga agcatgcaag tatgaaaccg atttgctcg gggtacgttt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
cacccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg aagcacgc tc atccgcctatg ccgaagagct tcttcggaaa 300  
aaggggcag accttttatg gtgcaacgcg aggacatctg cgagcgggtta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgaccggacc ccataattttg 420  
atgtataaga aattgacgta a 441

<210> 86

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 86

atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcattctc 60  
cgccgcatac agccgcttga agcatgcaag tatgaaaccg atttgctcg gggtgcgttt 120  
cacctcggtg gatattaccg gggcaagctg gtcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg aagcacgc tc atccgcctatg ccgaagagct tcttcggaaa 300  
aaggggcag accttttatg gtgcaacgcg aggacatctg cgagcgggtta ctataaaaag 360  
ctcggcttca gcbaacaggg caaagtctac gacataccgc cggtcggacc tcataattttg 420  
atgtataaga agttgacgta a 441

<210> 87

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 87

atgattgaag tcaaaccatat aaacgcggaa gatacgtatg agatcaggca ccgcattctc 60

cggccgaatc agccgctgga agcatgcaag tatgaaaccg atttgctcg gggtacgttt 120  
cacctcgcg gatattacccg gggcaagctg atcagcattcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actcgaagag 240  
taccgcgagc aaaaagcggg cagtcgctt atccgcccattg ccgaagagct tcttcggaga 300  
aaaggcgcgg acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc ctgtcggacc tcataattttg 420  
atgtataaga aattgacgta a 441

<210> 88  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 88  
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cggccgaatc agccgctgga agcatgcaag tatgaaaccg atttgctcg gggtacgttt 120  
cacctcggtg ggtactacccg gggcaagctg atcagcattcg cttcccttca taaagccgaa 180  
cattcagagc ttgagggcga agaacatgtat cagctgagag ggatggcgac gcttgaagga 240  
taccgtgagc aaaaagcggg cagtcgctt atccgctatg ccgaagagct tcttcgaaaa 300  
aaaggcgcgg acctttatg gtgcaacgcc aggacatctg tgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgatcggacc tcataattttg 420  
atgtataaga aattgacgta a 441

<210> 89  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 89  
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cggccgaatc agccgcttga agcatgtatg tatgaaaccg atttgctcg gggtacgctt 120  
cacctcggtg gatattacca gggcaagctg atcagcattcg cttcccttca taaagccgaa 180  
cattcaggccg ttgagggcga agaacatgtat cagctgagag ggatggcgac gctcgaaggg 240  
taccgcgagc aaaaagcggg cagtcgctt atccgcccattg ccgaagagct tcttcgaaaa 300  
aaaggcgcgg acctttatg gtgcaatgcc aggacatctg tgagcggcta ctatgaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgatcggacc tcataattttg 420  
atgtataaga aattgacgta a 441

<210> 90  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 90  
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cggccgaatc agccgctgga agcatgcaag tatgaaaccg atttgctcg gggcacgttt 120  
cacctcgcg gatattacca gggcaagctg atcagcattcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgcgagc aaaaagcggg aagtacgctc atccgcccattg ccgaagagct tcttcggaaaa 300  
aaggggcag acctttatg gtgcaatgcc aggacatctg tgagcgggta ctatgaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgatcggacc tcataattttg 420  
atgtataaga aattgacgta a 441

<210> 91  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 91  
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cgccgcgaaatc agccgctgga agcatgtatg tatgaaaaccg atttgctcg gggtgcgttt 120  
cacctcggtg gatattacca gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgcgagc aaaaagcggg aagtacgctt atccgcatcg ccgaagagct tcttcgaaaa 300  
aaaggcgcgg acctttatg gtgcaacgcc aggacatctg tgagcggta ctatgaaaag 360  
ctcggcttca gcaacaggg cgaagtctac gacataccgc cgatcggacc tcataattttg 420  
atgtataaga aattgacgta a 441

<210> 92  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 92  
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cgccgcgaaatc agccgctgga agcatgcaag tatgaaaaccg atttgctcg ggacgttt 120  
cacctcggtg gatattacccg gggcaagctg atcagcatcg cttcccttca taaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcaac gcttgaaggg 240  
taccgtgagc aaaaagcggg aagtacgctt atccgcatcg ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaacgcc aggacatctg cgagcggcta ctatgaaaag 360  
ctcggcttca gcaacaggg cgaagtctac gacataccgc cgatcggacc tcataattttg 420  
atgtataaga aattggcata a 441

<210> 93  
<211> 441  
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<220>  
<223> Synthetic DNA Sequence

<400> 93  
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cacctcggtg gatattacccg gggcaagctg atctgcacatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
taccgcgagc aaaaagcggg aagtacgctt atccgcatcg ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaatgcc aggacatctg tgagaggcta ctatgaaaag 360  
ctcggcttca gcaacaagg cgggtctac gatataccgc cgatcggacc tcataattttg 420  
atgtataaga aattggcgtta a 441

<210> 94  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 94

atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ctgcattctc 60  
cgccgaatc agccgcttga agcatgtatg tatgaaaccg atttgctcg gggtacgttt 120  
cacctcggtg gatattacca gggcaagctg atcagcatcg cttccttca taaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
taccgcgagc aaaaagcggg cagtacgctc atccgccatg ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaatgcc aggacatctg tgagcggcta ctataaaaag 360  
ctcggcttca gcbaacagg cgaaatctac gacataccgc cgatcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 95

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 95

atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcattctc 60  
cgccgaatc agccgcttga agcatgtatg tatgaaaccg atttgctcg gggtacgttt 120  
cacctcggtg gatattacca gggcaagctg atcagcatcg cttccttca taatgccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
taccgtgagc aaaaagcggg aagcacgctc atccgccatg ccgaagagct tcttcggaaa 300  
aaggggtag acctttatg gtgcaacgccc aggacatctg tgagcggcta ctataaaaag 360  
ctcggcttca gcbaacaagg cgggatctac gacataccgc cgatcggacc tcataatgg 420  
atgtataaga aattggcata a 441

<210> 96

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 96

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cacctcggtg, gatattaccc gggcaagctg atcagcatcg cttccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgcgagc aaaaagcggg aagtacgctt atccgccatg ccgaagagct tcttcggaaa 300  
aaaggcgcag acctttatg gtgcaacgccc aggacatctg tgagcggcta ctatgaaaag 360  
ctcggcttca gcbaacagg cgaaatctac gacataccgc cgatcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 97

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 97

atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcattctc 60  
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cacctcggtg gatattacca gggcaagctg atcagcatcg cttccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240

taccgtgagc aaaaagcggg aagcacgctc atccgccatg ccgaagagct tcttcggaaa 300  
aaaggcgca agccgttatg gtgcaacgcc aggacatctg tgagcggta ctataaaaag 360  
ctcggcttca gcaacaggg cgaagtctac gacataccgc cgatcggacc tcataaaaaag 420  
atgtataaga aattgacgta a 441

<210> 98  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 98  
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cacctcggtg gatattacca gggcaagctg atcagcatcg cttcccttca taaagccgaa 180  
cattcagagc ttgaaggcca aaaacagttt cagctgagag ggatggcgac gcttgaaggg 240  
taccgcgagc aaaaagcggg cagtacgctt atccgcctatg ccgaagagct tctccgaaaa 300  
aaaggcgccg acctttatg gtgcaatgcc aggacatctg cgagcggcta ctatgaaaaag 360  
ctcggcttca gcaacaggg cgaagtctac gacataccac cgatcggacc tcataaaaaag 420  
atgtataaga aattggcata a 441

<210> 99  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 99  
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cacctcggtg gatattatcg ggacaggctg atcagcatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagttt cagctgagag ggatggcgac gcttgaaggg 240  
taccgcgagc aaaaagcggg aagcacgctc atccgcctatg ccgaagagct tcttcggaaa 300  
aaaggcgccg acctttatg gtgcaacgcc aggacatctg tgagcggta ctataaaaag 360  
ctcggcttca gcaacaggg cgaagtctac gacataccgc cgatcggacc tcataaaaaag 420  
atgtataaga aactgacgta a 441

<210> 100  
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<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 100  
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cacctcggtg gatattaccc gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagttt cagctgagag ggatggcgac gcttgaaggg 240  
taccgcgagc aaaaagcggg aagtacgctt atccgcctatg ccgaagagct tcttcggaaa 300  
aaaggcgccg acctttatg gtgcaacgcc aggatctg tgagcggcta ctataaaaag 360  
ctcggcttca gcaacaagg cgggtctac gacataccgc cgatcggacc tcataaaaaag 420  
atgtataaga aattggcata a 441

<210> 101  
<211> 441

<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 101  
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catctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca taatgccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaagga 240  
taccgcgagc aaaaagcggg cagtaacgtt atccgcctg ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaatgcc aggacatctg cgagcggcta ctataaaaag 360  
ctcggcttca gcaacacagg cgaagtctac gacataccgc cgatcggacc tcataatgg 420  
atgtataaga aattggcata a 441

<210> 102  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 102  
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cgccgaatc agccgcttga agcatgtatg tatgaaaccg atttgctcg gggtacgttt 120  
cacctcggtg gatattatcg ggacaggctg atcagcatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
taccgcgagc aaaaagcggg cagtaacgtt atccgcctg ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaacgc aggacatctg cgagcggcta ctataaaaag 360  
ctcggcttca gcaacacagg cgaagtctac gacataccgc cgatcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 103  
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<220>  
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cacctcggtg gatattacca gggcaagctg atcagcaccg cttcccttca tcaagccgga 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgcgagc gaaaagcggg aagtacgctc atccgcctg ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaacgc aggatatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcaacaagg cgggtctac gatataccgc cgatcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

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<220>  
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<400> 104

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cacctcggtg gatattacca gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
taccgcgagc aaaaagcggg cagtcgctt atccgccatg ccgaagagct tcttcggaaa 300  
aaaggcgcag acctttatg gtgcaacgcc aggacatctg tgagcgggta ctataaaaag 360  
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<220>  
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cacctcggtg gatattaccc gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcggg cagtcgctt atccgccatg ccgaagagct tcttcggaaa 300  
aaggggcgcag acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
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atgtataaga aattgacgta a 441

<210> 106  
<211> 441  
<212> DNA  
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<220>  
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cacctcggtg gatattaccc gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
taccgcgagc aaaaagcggg cagtcgctt atccgccatg ccgaagagct tcttcggaaa 300  
aaaggcgcgg acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaagg cggggctac gacataccgc cgatcggacc ccataattttg 420  
atgtataaga aattgacgta a 441

<210> 107  
<211> 441  
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<220>  
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cacctcggtg gatattaccc gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgcgagc aaaaagcggg aagtacgctt atccgccatg ccgaagagct tcttcggaaa 300  
aaggggcgcag acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgatcggacc tcataattttg 420

atgtataaga aattgacgta a

441

<210> 108

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 108

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cacctcggtg gatattaccg gggcaagctg atcagcatcg cctcccttca tccagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagcggg cagtacgctt atccgccatg ccgaagagct tttcgaaaa 300  
aaaggcgcgg accctttatg gtgcaacgcc aggatatctg cgagcgggta ctatgaaaag 360  
ctcggcttca gcaaacaggg cgaagtctac gacataccgc cgatcggacc ccataatttg 420  
atgtataaga aattgacgta a 441

<210> 109

<211> 441

<212> DNA

<213> Artificial Sequence

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<223> Synthetic DNA Sequence

<400> 109

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cacctcggtg gatattaccg gggcaagctg atcagcatcg cctcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagcggg cagtacgctt atccgccatg ccgaagagct tttcgaaaa 300  
aaaggcgcgg accctttatg gtgcaacgcc aggacatctg cgagcggcta ctatgaaaag 360  
ctcggcttca gcaaacaggg cgaagtctac gacataccgc cgatcggacc tcataatttg 420  
atgtataaga aattgacgta a 441

<210> 110

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 110

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cacctcggtg gatattaccg gggcaagctg atcagcatcg cctcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagcggg aagtacgctt atccgccatg ccgaagagct tttcgaaaa 300  
aaggggcag accctttatg gtgcaacgcc aggatatctg cgagcggcta ctataaaaag 360  
ctcggcttca gcaacaagg cgggtctac gacataccgc cggtcggacc tcataatttg 420  
atgtataaga aattgacgta a 441

<210> 111

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 111

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cacctcggcg gatattaccg gggcaagctg atcagcatcg ctccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtggc aaaaagcggg cagtacgctt atccgcattg ccgaagagct tcttcggaaa 300  
aaaggcgcgg acctttatg gtgcaacgcg aggacatctg cgagcggcta ctataaaaag 360  
ctcggcttca gcaacaggg cgaagtctac gacataccgc cgatcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 112

<211> 441

<212> DNA

<213> Artificial Sequence

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<223> Synthetic DNA Sequence

<400> 112

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cacctcggcg gatattatca ggacaggctg atcagcatcg ctccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgcgagc aaaaagcggg aagtacgctt atccgcattg ccgaagagct tcttcggaaa 300  
aaaggcgcgg acctttatg gtgcaacgcg aggacatctg cgagcggcta ctataaaaag 360  
ctcggcttca gcaacaggg cgaagtctac gacataccgc cgatcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 113

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 113

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cacctcggcg gatattaccg gggcaagctg atcagcatcg ctccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gttgaaggg 240  
taccgcgagc aaaaagcggg cagtacgctt atccgcattg ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaacgcg aggacatctg cgagcggcta ctataaaaag 360  
ctcggcttca gcaacaggg cgaagtctac gacataccgc cgatcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 114

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 114

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cacctcggcg gatattatcg gggcaagctg atcagcatcg ctccttca tcaagccgaa 180

cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaagag 240  
taccgcgagc aaaaagcggg cagtacgctt atccgcccattt ccgaagagct tcttcggaaa 300  
aaaggcgcag accttttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgatcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 115  
<211> 441  
<212> DNA  
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<220>  
<223> Synthetic DNA Sequence

<400> 115  
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cacctcgggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg cagtacgctt atccgcccattt ccgaagagct tcttcggaaa 300  
aaaggcgcgg acctttgtg gtgcaacgcc aggacatctg cgagcgggta ctataaaag 360  
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atgtataaga aattgacgta a 441

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<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 116  
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ccataaaaagg tccgcgcctt tttccgaaag aagctcttcg gcatggcgaa tgagcgtgt 180  
tccccgtttt tgctcgggtt acccttcaag cgtccatc cctctcagct gatactgttt 240  
ttggccttca agctctgaat gttcggcttg atgaaaggag gcgatgctga tcagcttgcc 300  
ccggtaatataa ccaccggaggta gaaacgtgcc cccgagcaaa tcagtttcat acttgcattgc 360  
ttccagcggc tgattcggcc ggagaatgcg gtgcctgatc tcatacgtat cttccgcgtt 420  
tattggttt gttcaatca t 441

<210> 117  
<211> 441  
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<220>  
<223> Synthetic DNA Sequence

<400> 117  
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cacctcggcg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagg 240  
taccgcgagc aaaaagcggg cagtacgctt atccgcccattt ccgaagagct tcttcggaaa 300  
aaggggccag accttttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaagg cggggcttac gatataccgc cgatcggacc tcataatgg 420  
atgtataaga aattgacgta a 441

<210> 118

<211> 441  
<212> DNA  
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<220>  
<223> Synthetic DNA Sequence

<400> 118  
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cacctcggcg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgag 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaagag 240  
taccgcgagc aaaaagcggg cagtcgctt atccgcctg ccgaagagct tcttcggaaa 300  
aaggggcag accttttatg gtgcaacgcc aggacatctg cgagcggcta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgatcggacc tcataatttg 420  
atgtataaga aattgacata a 441

<210> 119  
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<220>  
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<400> 119  
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cacctcggcg gatattaccg gggcaagctg atcagcatcg cttcccttca taatgccgaa 180  
cattcagagc ttgatggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgcgagc aaaaagcggg aagcacgctc atccgcctg ccgaagagct tcttcggaaa 300  
aagggcgcag accttttatg gtgcaacgcc aggacatctg tgagcggcta ctataaaaag 360  
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atgtataaga aattgacgta a 441

<210> 120  
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<220>  
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cacctcggcg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
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taccgcgagc aaaaagcggg aagcacgctc atccgcctg ccgaagagct tcttcggaaa 300  
aaggggcag accttttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
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atgtataaga aattggcata a 441

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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttccccc 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
taccgcgagc taaaagcggg aagtacgctt atccgc 300  
aaaggcgcgg acctttatg gtgcaacgccc 360  
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atgtataaga aattgacgta a 441

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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttccccc 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg aagcacgctc atccgc 300  
aaaggcgcgg acctttatg gtgcaacgccc 360  
ctcggttca gcaacaagg cgggtctac gatataccgc cgatcggacc tcataaaaaag 420  
atgtataaga aattgacgta a 441

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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttccccc 180  
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aaggggcag acctttatg gtgcaacgccc 360  
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atgtataaga aattgacata a 441

<210> 124  
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<212> DNA  
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<220>  
<223> Synthetic DNA Sequence

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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttccccc 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
taccgcgagc aaaaagcggg aagcacgctc atccgc 300  
aaaggcgcgg acctttatg gtgcaacgccc 360

ctcggcttca gctaaccaggcgaa gatacgtatg agatcaggca ccgcatttc 420  
atgtataaga aattgacgta a 441

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<220>  
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cacctcggtg 120  
gatattaccg gggcaagctg atcagcatcg cttccccc 180  
tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac 240  
taccgcgagc 240  
aaaaagcggg tagtacgctt atccgc 300  
ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaacgccc 360  
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atgtataaga aattgacgta a 441

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cacctcggtg 120  
gatattaccg gggcaagctg atcagcatcg cttccccc 180  
tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac 240  
taccgcgagc 240  
aaaaagcggg cagtacgctt atccgc 300  
ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaacgccc 360  
aggacatctg tgagcgggta ctataaaaag 360  
ctcggcttca gctaaccaggcgaa gatacgtatg cgaagtctac 420  
gacataccgc cgatcggacc tcataattttg 420  
atgtataaga aattgacgta a 441

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<220>  
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cacctcggtg 120  
gatattaccg gggcaagctg atcagcatcg cttccccc 180  
tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac 240  
taccgtgagc 240  
aaaaagcggg cagtacgctt atccgc 300  
ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaacgccc 360  
aggacatctg cgagcggcta ctataaaaag 360  
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gacataccgc cgatcggacc tcataattttg 420  
atgtataaga aattaacata a 441

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cacttaggcg gctttacag gggcaagctg atttccatag cttcattcca ccaggccgag 180  
catccagaac tccaggggca gaaaacaatac caactccgag gtatggctac cttggaaggt 240  
tatcgtgacc agaaaagcggg atcgagccta attaaacacg ctgaacagat cttcggaaag 300  
cggggggcgg acatgctatg gtgcaatgcg cggacatccg ccgctggcta ctacaaaaaag 360  
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atgtataaac gcctcacata a 441

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<220>  
<223> Synthetic DNA Sequence

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cacctcggtg gatattaccg aggcaagctg atcagcatcg cttccattcca tcaagccgaa 180  
cattcagagc ttgaaggcca taaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcgagc aaaaagcggg aagcacgcctc atccgcctatg ccgaagagct tttcggaaa 300  
aagggggcag acctttatg gtgcaacgcg aggacatctg cgagcggctt ctataaaaaag 360  
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atgtataaga aattgacgta a 441

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cacctcggtg gatattaccg aggcaagctg atcagcatcg cttccattcca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
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aaaggcgcgg acctttatg gtgcaacgcg aggacatctg cgagcgggtt ctataaaaaag 360  
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cacctcggtg gatattaccg gggcaagctg atcagcatcg ctccttca ccaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagga 240  
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aagggggcag acctttatg gtgcaacgcc aggacatctg cgagcggta ctataaaaag 360  
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aagggggcag acctttatg gtgcaacgcc aggacatctg cgagcggcta ctataaaaag 360  
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aagggggcag acctttatg gtgcaacgcc aggacatctg cgagcggta ctataaaaag 360  
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cacctcggtg gatattatcg gggcaagctg atcagcatcg cttccattca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaagga 240  
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aagggggcag acctttatg gtgcaacgcc aggatatctg cgagcggta ctataaaaag 360  
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atgtataaga aattgacgta a 441

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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgcgagc aaaaagcggg aagcacgctc atccgcatg ccgaagagct tcttcggaaa 300  
aaaggcgcgg acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
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atgtataaga aattgacgta a 441

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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
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aaggggcag acctttatg gtgcaacgcc aggacatctg cgagcgggta ctataaaaag 360  
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cacctcggtg gatattatcg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
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atgtataaga aattgacgta a 441

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atgtataagg aattgacgta a 441

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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
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aaaggcgcgg accttttatg gtgcaacgcc aggacatctg cgagcggta ctataaaaag 360  
ctcggttca gcgaacaagg cgggtctac gacataccgc cggtcggacc tcataatttg 420  
atgtataaga aattgacgta a 441

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cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagga 240  
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aaaggcgcgg accttttatg gtgcaacgcc aggacatctg cgagcggcta ctataaaaag 360  
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atgtataaga aattgacgta a 441

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atgtataaga aattgacgta a 441

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<210> 146

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atgtataaga aattgacgta a 441

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atgtataaga aattgacata a 441

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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagga 240  
taccgtgagc aaaaagcggg cagtacgctt atccgccc 300  
aaaggcgcgg acctttgtg gtgcaacgccc aggacgtctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgaccggacc ccataaaaa 420  
atgtataaga aattgacgta a 441

<210> 180  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 180  
atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgcgatc agccgctgga agcatgcaag tatgaaaactg atttgcgtgg 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcggg cagtacgctt atccgccc 300  
aaggcgcgg accttatatg gtgcaacgccc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cggtcggacc tcataaaaa 420  
atgtataaga aattgacata a 441

<210> 181  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 181  
atgattgaag tcaaaccaat aaatgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgcgatc agccgcttga agcatgtatg tatgaaaaccg atttgcgtcg 120  
caccttaggtg gatattaccg gggcaagctg atcagcatcg cttccctttaa tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcggg cagtacgctt atccgccc 300  
aaggggcag acctttatg gtgcaacgccc aggacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cggtcggacc tcataaaaa 420  
atgtataaga aattgacata a 441

<210> 182  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 182  
atgattgaag tcaaaccaat aaccgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgttga agcatgcaag tatgaaacccg atttgctcg gggtacgtt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cctccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac actagaaggg 240  
taccgcggc aaaaagcggg cagtaacgctc atccgcattg ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaacgccc agaacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggc cgaagtctac gacataccgc cgaccggacc ccatattttg 420  
atgtataaga aattgacgta a 441

<210> 183  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 183  
atgattgaag tcaaaccaat aaacgcggaa gatgcgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgttga agcatgcaag tatgaaacccg atttgctcg gggcacgtt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cctccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaagag 240  
taccgcggc aaaaagcggg aagtacgctt atccgcattg ccgaagagct tcttcggaaa 300  
aaggggcgg acctttatg gtgcaacgccc agaacatctg cgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggc cgaagtctac gacataccgc ctgtcggacc tcataattttg 420  
atgtataaga aattgacgta a 441

<210> 184  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 184  
atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgttga agcatgcaag tatgaaacccg atttgctcg gggtacgtt 120  
cacctcggtg gatattaccg gggcaagctg atctgcattg cctccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
taccgcggc aaaaagcggg aagtacgctt atccgcattg ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaatgcc agaacatctg tgagaggcta ctatgaaaag 360  
ctcggcttca gcbaacaagg cggggctac gatataccgc cgatcggacc tcataattttg 420  
atgtataaga aattggcgta a 441

<210> 185  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 185  
atgattgaag tcaaaccaat aaacgcggaa gatacgtatg agatcaggca ccgcatttc 60  
cgccgaatc agccgttga agcatgcaag tatgaaacccg atttgctcg gggtacgtt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttccttca taaagccgaa 180  
cattcagagc ttgagggcgta agaacatgt cagctgagag ggatggcgac gcttgaaggg 240  
taccgtgagc aaaaagcggg aagcacgctc atccgcattg ccgaagagct tcttcggaaa 300  
aaggggcag acctttatg gtgcaatgcc agaacatctg tgagcgggta ctataaaaag 360

ctcggcttca gccaacaggcg aactaaggca taaaatactc 420  
atgtataaga aattgacgta a 441

<210> 186  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 186  
atgatagaag taaaaccgt taacgcagag gatacctatg aactaaggca taaaatactc 60  
agaccaaacc agccgataga agcgtgtatg tatgaaagcg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tcgaaggcca gaaacagtac cagctccgag gtatggctac cttggaaagg 240  
tatcgtgatc agaaaagcggg atcgactcta attaaacacg ctgaagaaat tcttcgttaag 300  
agggggcgg acatgctttg gtgcaatgcg cggacaaccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gatacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 187  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 187  
atgatagagg taaaaccgt taacgcagag gatacctatg aactaaggca taaaatactc 60  
agaccaaacc agccgataga agcgtgtatg tatgaaagcg atttacttcg tgggcattt 120  
cacttaggcg gctttacgg gggcaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tcgaaggcca gaaacagtac cagctccgag gtatggctac cttggaaagg 240  
tatcgtgatc agaaaagcggg atcgactcta attaaacacg ctgaacaact tcttcgttaag 300  
agggggcgg acatgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gaaacgcccgc cagtaggacc tcacatcctg 420  
atgtataaaa agatcaca 438

<210> 188  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 188  
atgctagagg taaaaccgt taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgataga agcgtgtatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaactg atttccatag cttcattcca ccaggccgag 180  
cactcagaac tcgaaggcca gaaacagtac cagctccgag gtatggctac cttggaaagg 240  
tatcgtgatc agaaaagcggg atcgagtcta attaaacacg ctgaacaact tcttcgttaag 300  
agggggcgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gatacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcaca 438

<210> 189  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 189  
atgcttagagg taaaactgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgttata agcgtgtatg tatgaaaccc atttacttcg tggtgcattt 120  
cacttaggcg gctttacag gggcaactg atttccatag cttcattcca ccaggcccag 180  
cactcagacc tcgaaggcca gaaacagtac cagctccgag gtatggctac ctttggaaagg 240  
ttcgtgtatc agaaaaggccc atcgagtcta attaaacacg ctgaagaaat tcttcgttaag 300  
aggggggcga acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gatacgcgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcaca 438

<210> 190  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 190  
atgatagagg taaaacctgat taacgcagag gatacctatg aactaaggca taaaataactc 60  
agaccaaacc agccgtataga agcgtgtatg tatgaaaccc atttacttcg tggtgcattt 120  
cacttaggcg gctttacag gggcaactg atttccatag cttcattcca ccaggcccag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac ctttggaaagg 240  
tatcgtgtatc agaaaaggccc atcgagtcta attagacacg ctgaacaaaat tcttcgttaag 300  
aggggggcgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gatacgcgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcaca 438

<210> 191  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 191  
atgatagagg taaaacctgat taacgcagag gatacctatg aactaaggca taaaataactc 60  
agaccaaacc agccgttata agcgtgtatg tatgaaaccc atttacttcg tggtgcattt 120  
cacttaggcg gctttacag gggcaactg atttccatag cttcattcca ccaggcccag 180  
cactcaggcc tccaaggcca gaaacagtac cagctccgag gtatggctac ctttggaaagg 240  
tatcgtgtatc agaaaaggccc atcgagtata attaaacacg ctgaagaaat tcttcgttaag 300  
aagggggcgg acttgctttg gtgcaatgcg cggacgtccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gacacgcgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcaca 438

<210> 192  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 192  
atgatagagg taaaacctgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgtataga agcgtgtatg tatgaaaccc atttacttcg tggtgcattt 120

cacttaggcg gctttacag gggcaaactg attccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac ctttgcagg 240  
tatcgtgagc agaaaagcggg atcgactcta attagacacg ctgaacaact tttcgtaag 300  
agggggggcgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaagg 360  
ttaggcttca gcgagcaggg agaggtat tt gatacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcaca 438

<210> 193  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 193  
atgcttagagg tgaaaccgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgataga agcgtgtatg tatgaaagcg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaaactg attccatag cttcattcca ccaggccgag 180  
cactcagaaac tccaaggcca gaaacagtac cagctccgag gtatggctac ctttgcagg 240  
tatcgtgagc agaaaagcggg atcgactcta attaaacacg ctgaagaaat tttcgtaag 300  
agggggggcgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaagg 360  
ttaggcttca gcgagcaggg agagatattt gatacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcaca a 441

<210> 194  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 194  
atgatagaag tgaaaccgat taacgcagag gagacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgataga agcgtgtatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaaactg attccatag cttcattcca ccaggccgag 180  
cactcagaaac tcgaaggcca gaaacagtac cagctccgag gtatggctac ctttgcagg 240  
tatcgtgatc agaaaagcggg atcgactcta attagacacg ctgaacaact tttcgtaag 300  
aagggggcga atatgctttg gtgcaatgcg cggacaaccg cctcaggcta ctacaaaagg 360  
ttaggcttca gcgagcaggg agagatattt gatacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcaca 438

<210> 195  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 195  
atgcttagagg tgaaaccgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgataga agcgtgtatg tatgaaagcg atttacttcg tgggcactt 120  
cacttaggcg gctttacag gggcaaactg attccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac ctttgcagg 240  
tttcgtgatc agaaaagcggg atcgactcta attagacacg ctgaacaaat tttcgtaag 300  
agggggggcgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaagg 360  
ttaggcttca gcgagcaggg aaaggtat tt gatacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcaca 438

<210> 196  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 196  
atgcttagagg taaaaccgat taacgcagag gatacctatg aactaaggca taaaataactc 60  
agaccaaacc agccgttaga agtgttatg tatgaaaccg atttacttcg tggtgcattt 120  
cacttaggcg gctttacag gggcaactg atttccatag cttcattcca ccaggcccag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac ctttgcgtt 240  
tatcgatc agaaaagcggg atcgagtcta attagacacg ctgaacaaat tcttcgttaag 300  
agggggcgg acatgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gaaacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcaca 438

<210> 197  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 197  
atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcatttctc 60  
cgccccgaaatc agccgcttgg a gcatgcaag tatgaaaccg atttgcgtt gggcacgcgtt 120  
cacctcggtg gatattaccg gggcaagctg gtcagcattcg cttccatttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggtatggcgac gcttgaaggg 240  
taccgtgagc aaaaagcggg cagtacgtt atccgcatttgc ccgaagagct tcttcggaaa 300  
aaggggcgg acctttatg gtgcaacgcg aggacatctg cgagcgggta ctataaaaaag 360  
ctcggttca gcgAACAGGG CGAAGTCTAC GACATACCGC CGGTGGGACC TCAATTTCG 420  
atgtataaga aattgacgta a 441

<210> 198  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 198  
atgatagagg taaaaccgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgtatg agcgttatg tatgaaagcg atttacttcg tggtgcattt 120  
cacttaggcg gcttattacag gggcaactg atttccatag cttcattcca ccaggcccag 180  
cactcagaac tccaaggcca gaaacagtac cagctccgag gtatggctac ctttgcgtt 240  
tttcgtgagc agaaaagcggg atcgactcta attagacacg ctgaacaaat tcttcgttaag 300  
agggggcgg acatgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gatacgcgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcaca 438

<210> 199  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 199  
atgatagagg taaaaccgt taacgcagag gatacctatg aactaaggca taaaatactc 60  
agaccaaacc agccgataga agcgtgtatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctttacgg gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tcgaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcggg atcgactcta attagacacg ctgaagaaat tcttcgttaag 300  
aagggggcga acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gatacgcgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcaca 438

<210> 200  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<221> misc\_feature  
<222> 134, 313  
<223> n = A,T,C or G

<400> 200  
atgatagagg taaaaccgt taacgcagag gatacctatg aactaaggca taaaatactc 60  
agaccaaacc agccgttaga agtgtgtatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctnttacag gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagaac tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgatc agaaaagcggg atcgagtcta attaaacacg ctgaacaaat tcttcgttaag 300  
agggggcgg acntgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gatacgcgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 201  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 201  
atgatagagg taaaaccgt taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgataga agtgtgtatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcggg atcgactcta attagacacg ctgaacaaat tcttcgttaag 300  
agggggcgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gatacgcgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 202  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 202  
atgcttagagg taaaaccgt taacgcagag gatacctatg aactaaggca tagaataactc 60

agaccaaacc agccgataga agcgtgtatg tttgaaagcg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaactg atttccatag cttcattcca ccaggccgag 180  
cactcagaac tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcggg atcgagtcta attagacacg ctgaagaaat tcttcgttaag 300  
agggggccgg acatgctttg gtgtaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gatacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 203  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 203  
atgatagagg taaaaccgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgataga agtgtgtatg tatgaaaccc atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaactg atttccatag cttcattcca ccaggccgag 180  
cactcagaac tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
ttcgtgagc agaaaagcggg atcgagtcta attagacacg ctgaacaaaat tcttcgttaag 300  
agggggccgg acttgctttg gtgtaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gatacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 204  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 204  
atgatagagg taaaaccgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgataga agcgtgtatg tttgaaagcg atttacttcg tgggcattt 120  
cacttaggcg gctattacag gggcaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
taccgcgatc agaaaagcggg atcgagtcta attagacacg ctgaacaaaat tcttcgttaag 300  
agggggccgg acttgctttg gtgtaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gatacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 205  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 205  
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cacttaggcg gctttacag gggcaactg atttccatag cttcattcca ccaggccgag 180  
cactcagaac tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcggg atcgagtcta attaaacacg ctgaagaaat tcttcgttaag 300  
agggggccgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gatacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 206  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 206  
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agaccaaacc agccgataga agtgttatg tatgaaagcg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaaaactg atttccatag cgtcattcca ccaggccgag 180  
caccaggacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgatc agaaaagcggg atcgagtcta attaaacacg ctgaacaaat tcttcgttaag 300  
agggggccgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gaaacgcgc cagtagggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 207  
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<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 207  
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cacttaggcg gctttacag gggcaaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac caactccgag gtatggctac cttggaaggt 240  
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agggggccgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaagg 360  
ttaggcttca gcgagcaggg agagatattt gacacgcgc cagtagggcc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 208  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 208  
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cacttaggcg gctttacag gggcaaaactg atttccatag cgtcattcca ccaggccgag 180  
cactcggAAC tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgatc agaaaagcggg atcgagtcta attagacacg ctgaacaaat tcttcgttaag 300  
agggggccgg acatgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gatacgcgc cagtagggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 209  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 209

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cacttggcg gctttacgg gggcaaactg atttccatag cgtcattcca ccaggccgag 180  
caccaggacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgatc agaaaagcggg atcgagtcta attagacacg ctgaacaact tcttcgtaag 300  
agggggcgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattc gaaacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 210

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 210

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cacttaggcg gctttacag gggcaaactg atttccatag cgtcattcca ccaggccgag 180  
cactcagaac tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcggg atcgagtcta attaaacacg ctgaagaaat tcttcgtaag 300  
agggggcgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gacacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 211

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 211

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cacttaggcg gctttacgg gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcggg atcgactcta attagacacg ctgaagaaat tcttcgtaag 300  
agggggcgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gatacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 212

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 212

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cacttaggcg gctttacag gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240

tatcgtgagc agaaagcggg atcgagtcta attaaacacg ctgaacaaat tcttcgtaag 300  
agggggcgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gatacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 213  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 213  
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agaccaaacc agccgataga agtgttatg tatgaaaccc atttacttcg tggcattt 120  
cacttaggcg gctttacgg gggcaactg atttccatag cttcattcca ccaggccgag 180  
cactcagaac tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaagg 240  
tatcgtgatc agaaagcggg atcgagtcta attaaacacg ctgaagaaat tcttcgtaag 300  
agggggcgg acttgctttg gtgtaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gaaacgcccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 214  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 214  
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cgccctaattt agccgataga ggcatgcattg tatgaaagcg atctgctgcg gggctcggttc 120  
catggggcg ggttctatcg tggccaaattt atctcgattt cgagtttcca caaagctgaa 180  
cactcagaac tgcaggcga aaagcagtat caattacgag ggatggcgcac cctcgaagga 240  
ttccgtgagc agaaggctgg ctcttcgctt attaggcacg ccggaggagat actacggaaat 300  
aaaggccag atctgctttg gtgtaatgca cgcacgacag cctccggta ctataaaagg 360  
cttggttta gtgagcacgg cgaaggttt gaaacccccgc cgggttgggcc gcacattctt 420  
atgtacaaaaa gaatcact 438

<210> 215  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 215  
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cgccctaattt agccgtttaga ggcatgcattg tatgaaagcg atctgctgcg gggctcggttc 120  
catggggcg ggttctatcg tggccaaattt atctcgattt cgagtttcca ccaagctgaa 180  
cactcagaac tggaggcga aaagcagtat caattacgag ggatggcgcac cctcgaagga 240  
ttccgtgagc agaaggctgg ctcttcgctt attaggcacg ccggaggagat actacggaaa 300  
agggggcag atctgctttg gtgtaatgca cgcacgacag cccgggtta ctataaaaagg 360  
cttggttta gtgagcacgg cgaaggttt gacacccccgc cgggttgggcc gcacattctt 420  
atgtacaaaaa gaatcact 438

<210> 216  
<211> 438

<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 216  
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cgccctaatac agccgttaga ggcatgcattg tatgaaagcg atctgctgcg gggctcggttc 120  
catggggcg ggttctatcg tggcaaaatttgcg atctcgatttgcg 180  
caattacgag cgatggcgac cctcgaagga 240  
taccgtgatc agaaggctgg ctctcgctt attaggcacttgcg 300  
agaggggcag atctgctttgcg 360  
cttggttta gtgagcaggg cgaagtttc gacacccccgcg 420  
atgtacaaaaa aactcact 438

<210> 217  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 217  
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cgccctaatac agccgttaga ggcatgcattg tatgaaagcg atctgctgcg gggctcggttc 120  
catggggcg ggttctatcg tggcaaaatttgcg atctcgatttgcg 180  
caattacgag cgatggcgac cctcgaagga 240  
taccgtgatc agaaggctgg ctctacgttgcg attaaggcacttgcg 300  
aaaggggcag atctgctttgcg 360  
cttggttta gtgagcaggg cgaattttc gacacccccgcg 420  
atgtacaaaaa gactcact 438

<210> 218  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 218  
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catggggcg ggttctatcg tggcaaaatttgcg atctcgatttgcg 180  
caattacgag cgatggcgac cctcgaagga 240  
ttccgtgatc agaaggctgg ctctcgctt attaaggcacttgcg 300  
agaggggcag atctgctttgcg 360  
cttggttta gtgagcaggg cgaattttc gaaacccccgcg 420  
atgtacaaaaa gactcact 438

<210> 219  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 219

atgctagaag taaaacccat taacgcagag gagacttacg aacttcgaca caagatcctg 60  
cgccctaatac agccgataga ggcatgcatg tatgaaaccg atctgctgcg gggctcggtc 120  
catttggcg ggttctatcg tggcaattt atctcgattt cgagtttcca ccaagctgaa 180  
cactcagacc tgcaaggcgca aaagcagtat caattacgag ggatggcgac cctcgaagga 240  
taccgtgagc agaaggctgg ctctacgctt attaaggcact cccgaggagct actacggaaa 300  
aaagggcgag atatgctttt gtgcaatgca cgcacgacag ccgcgggtta ctataaaaagg 360  
cttgggttta gtgagcaggg cgaagtttc gacaccccgcc cggttggccc gcacatttt 420  
atgtacaaaaaa aaatcact 438

<210> 220  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 220  
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cgccctaatac agccgttaga ggcatgcatg tatgaaaccg atctgctgcg gagcgcattc 120  
catttggcg ggttctatcg tggcaattt atctcgattt cgagtttcca caaagctgaa 180  
cactcagaac tgcaaggcgca aaagcagtat caattacgag ggatggcgac cctcgaagga 240  
taccgtgatc agaaggctgg ctcttcgctt attaggcact cccgaggagat actacggaaa 300  
agagggcgag atatgctttt gtgcaatgca cgcacgtcag ccgcgggtta ctataaaaagg 360  
cttgggttta gtgagcaggg cgaagtttc gacaccccgcc cggttggccc gcacatttt 420  
atgtacaaaaaa gaatcactta a 441

<210> 221  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 221  
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cgccctaatac agccgataga ggcatgcatg tatgaaaccg atctgctgcg gggctcggtc 120  
catttggcg ggttctatcg tggcaattt atctcgattt cgagtttcca ccaagctgaa 180  
cactcagacc tgcaaggcgca aaagcagtat caattacgag ggatggcgac cctcgaagga 240  
taccgtgagc agaaggctgg ctcttcgctt attaaggcact cccgagcagct actacggaaa 300  
aaagggcgag atatgctttt gtgtaatgca cgcacgtcag ccgcgggtta ctataaaaagg 360  
cttgggttta gtgagcacgg cgaattttc gaaaccccgcc cggttggccc gcacatttt 420  
atgtacaaaaaa gaatcact 438

<210> 222  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 222  
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cgccctaatac agccgttaga ggcatgcatg tatgaaaccg atctgctgcg gggctcggtc 120  
catttggcg ggttctatcg tggcaattt atctcgattt cgagtttcca ccaagctgaa 180  
cattcagaac tggaaaggcgca aaagcagtat caattacgag ggatggcgac ttcgaagga 240  
taccgtgatc agaaggctgg ctcttcgctt attaggcact cccgaggagat actacggaaa 300  
agagggcgag atatgctttt gtgcaatgca cgcacgacag ccgcgggtta ctataaaaagg 360  
cttgggttta gtgagcaggg cgaattttac gacaccccgcc cggttggccc gcacatttt 420

atgtacaaaa aactcact

438

<210> 223

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 223

atgatagaag taaaacccat taacgcagag gagacttacg aacttcgaca caagatcctg 60  
cgccctaattc agccgttaga ggcatgcatt tatgaaaccg atctgctgcg gggcgcggttc 120  
catggggcg ggttctatcg tggcaaattt atctcgattt cgagtttcca ccaagctgac 180  
cactcagaac tgcaaggggca aaaggcgttat caattacgag ggatggcgac cctcgaagga 240  
taccgtgagc agaaggctgg ctctacgctt attaggcactt ccgagcagat actacggaaa 300  
agaggggcag atactactttt gtgcaatgca cgcacgttag cggccgggtta ctataaaaag 360  
cttgggtttta gtgagcacgg cgaaattttc gaaaccccgcc cggttgggccc gcacattttt 420  
atgtacaaaa gactcactta a 441

<210> 224

<211> 438

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 224

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cgccctaattc agccgttaga ggcatgcatt tatgaaagcg atctgctgcg gggcgcggttc 120  
catggggcg ggttctatcg tggcaaattt atctcgattt cgagtttcca ccaagctgaa 180  
cactcagacc tgcaaggggca aaaggcgttat caattacgag ggatggcgac cctcgaagga 240  
taccgtgagc agaaggctgg ctcttcgctt attaggcactt ccgaggagat actacggaaa 300  
aaaggggcag atatgctttt gtgcaatgca cgcacgacag cggccgggtta ctataaaagg 360  
cttgggtttta gtgagcacggg cgaaattttc gacaccccgcc cggttgggccc gcacattttt 420  
atgtacaaaa gaatcact 438

<210> 225

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 225

atgatagagg taaaaccggat taacgcagag gatacctatg aactaaggca tagaataactc 60  
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cacttaggcg gcttttacag gggcaaactg atttccatag cttcatttcca ccaggccgag 180  
cactcagacc tccaaaggcca gaaacagtac cagctccgag gtatggctac ttggaaaggt 240  
tatcgtgagc agaaaaggccc atcgagtctt attaaacacg ctgaacaaat tcttcgttaag 300  
agggggccgg acttgctttt gtgcaatgca cggacatccg cctcaggctt ctacaaaaag 360  
ttaggcttca gcgagcacggg agaggtattt gatacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 226

<211> 438

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 226

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catttggcg ggttctatcg tggccaaatttgcg atctcgatttgcg ccaagctgaa 180  
cactcagaac tgcaagggca aaagcagtat caattacgag ggatggcgac cctcgaagga 240  
taccgtgagc agaaggctgg ctctacgctt attaaggcacttgcg cccgaggagat actacggaaa 300  
aaaggggcag atatgctttg gtgcaatgca cgcacgtcag cccgcccgtta ctataaaagg 360  
cttgggttttgcg atatgctttg gtgcaatgca cgcacgtcag cccgcccgtta ctataaaagg 420  
atgtacaaaaa gaatcact 438

<210> 227

<211> 438

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 227

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catttggcg ggtactatcg tggccaaatttgcg atctcgatttgcg caaagctgaa 180  
cactcagaac tgcaagggca aaagcagtat caattacgag ggatggcgac cctcgaagga 240  
taccgtgagc agaaggctgg ctctacgctt attaaggcacttgcg cccgaggagat actacggaaa 300  
aaaggggcag atatgctttg gtgcaatgca cgcacgtcag cccgcccgtta ctataaaagg 360  
cttgggttttgcg atatgctttg gtgcaatgca cgcacgtcag cccgcccgtta ctataaaagg 420  
atgtacaaaaa aactcact 438

<210> 228

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 228

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catttggcg ggttctatcg tggccaaatttgcg atctcgatttgcg caaagctgaa 180  
cactcagacc tggaagggca aaaccagtat caattacgag ggatggcgac cctcgaagga 240  
taccgtgagc agaaggctgg ctctacgctt attaggcacttgcg cccgaggagat actacggaaa 300  
agaggggcag atatgctttg gtgcaatgca cgcacgtcag cccgcccgtta ctataaaagg 360  
cttgggttttgcg atatgctttg gtgcaatgca cgcacgtcag cccgcccgtta ctataaaagg 420  
atgtacaaaaa gactcactta a 441

<210> 229

<211> 438

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 229

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catttggcg ggttctatcg tggccaaatttgcg atctcgatttgcg caaagctgaa 180

cactcagacc tgaaaggca aaaggcgtat caattacgag ggatggcgac cctcgaagga 240  
taccgtgagc agaaggctgg ctctacgctt attaggcacg ccgagcagat actacggaaa 300  
agaggggcag atatgcctg gtgcaatgca cgcacgtcag ccgcccgtta ctataaaagg 360  
cttggttta gtgagcaggg cgaagtttc gaaaccccgc cggttggcc gcacatttt 420  
atgtacaaaaa gactcact 438

<210> 230  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 230  
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catggggcg ggttctatcg tggcaaatttgcg atctcgatttgcg ccaagctgaa 180  
cactcagacc tgcaaggca aaaggcgtat caattacgag ggatggcgac cctcgaagga 240  
taccgtgagc agaaggctgg ctctacgctt attaggcacg ccgagcagct actacggaaa 300  
agaggggcag atatgccttgcg atatgccttgcg ccgcccgtta ctataaaagg 360  
cttggttta gtgagcacgg cgaagtttc gacaccccgccggttggcc gcacatttt 420  
atgtacaaaaa gactcact 438

<210> 231  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 231  
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catggggcg ggtactatcg tggccaaatttgcg atctcgatttgcg ccaagctgaa 180  
cactcagaac tgcaaggca aaaggcgtat caattacgag ggatggcgac cctcgaagga 240  
ttccgtgagc agaaggctgg ctctacgctt attaaggcacg ccgagcagat actacggaaa 300  
agaggggcag atatgccttgcg atatgccttgcg ccgcccgtta ctataaaagg 360  
cttggttta gtgagcacgg cgaatatttc gacaccccgccggttggcc gcacatttt 420  
atgtacaaaaa aactcactta a 441

<210> 232  
<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 232  
atgatagaag tgaaacctat taacgcagag gagacttacg aacttcgaca caggatcctg 60  
cgccctaatac agccgttaga ggcatgcattg tatgaaagcg atctgctgcg gggctcggttc 120  
catggggcg ggttctatcg tggccaaatttgcg atctcgatttgcg ccaagctgaa 180  
cactcagacc tagaaggca aaaggcgtat caattacgag ggatggcgac cctcgaagga 240  
taccgtgatc agaaggctgg ctctacgctt attaaggcacg ccgaggagct actacggaaa 300  
agaggggcag atatgccttgcg atatgccttgcg ccgcccgtta ctataaaagg 360  
cttggttta gtgagcacgg cgaatatttc gaaaccccgccggttggcc gcacatttt 420  
atgtacaaaaa aaatcact 438

<210> 233

<211> 438  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 233  
atgatagaag tgaaaacctat taacgcagag gatacttacg aacttcgaca caagatcctg 60  
cgccctaatac agccgataga ggcatgcatt tatgaaagcg atctgctgcg gggctcggtc 120  
catttggcg ggttctatcg tggccaattt atctcgattt cgagtttcca ccaagctgaa 180  
cactcagacc tggaaaggcga aaaggcagtat caattacgag ggatggcgac cctcgaagga 240  
taccgtgatc agaaggctgg ctttcgctt attaagcacg ccgaggagat actacggaaa 300  
agaggggcag atctgctttg gtgcaatgca cgcacgtcag ccgcccgtta ctataaaaagg 360  
cttggttta gtgagcaggg cgaaatttc gacacccccgc cggttgggcc gcacatttt 420  
atgtataaaaa aaatcact 438

<210> 234  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 234  
atgcttagagg tgaaaaccgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgataga agcgtgtatg tatgaaagcg atttacttcg tgggcattt 120  
cacttaggcg gctttagtacag gggcaaactg atttccatag cttcatttcca ccaggccgag 180  
cactcagaac tccaaaggcca gaaacagtac cagctccgag gtatggctac ctttggaaagg 240  
tatcgtgagc agaaaaggcgg atcgagtctt tttaaacacg ctgaagaaaat tttcgtaag 300  
agggggggcgg acttgctttg gtgtaatgca cggacatccg cctcaggctt ctacaaaaagg 360  
ttaggcttca gcgagcaggg agagatattt gaaacgcccgc cagtaggacc tcacatcctg 420  
atgtataaaaa ggatcacata a 441

<210> 235  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 235  
atgcttagagg tgaaaaccgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgataga agcgtgtatg tatgaaagcg atttacttcg tgggcattt 120  
cacttaggcg gcttttagtacag gggcaaactg atttccatag cttcatttcca ccaggccgag 180  
cactcagaac tccaaaggcca gaaacagtac cagctccgag gtatggctac ctttggaaagg 240  
tatcgtgagc agaaaaggcgg atcgagtctt tttaaacacg ctgaagaaaat tttcgtaag 300  
agggggggcgg acttgctttg gtgtaatgca cggacatccg cctcaggctt ctacaaaaagg 360  
ttaggcttca gcgagcaggg agagatattt gaaacgcccgc cagtaggacc tcacatcctg 420  
atgtataaaaa ggctcacata a 441

<210> 236  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 236  
atgcttagagg taaaaccgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgttata agcgtgtatg tatgaaagcg atttacttcg tggtgcattt 120  
cacttaggcg gctattacgg gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagaac tccaaggcca gaaacagtac cagctccgag gtatggctac ctttgcgttaag 240  
tatcgtgagc agaaaagcggg atcgagtcta attaaacacg ctgaagaaat tttcgtaag 300  
agggggccgg acttgctttg gtgtaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gaaacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 237  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 237  
atgatagaag taaaaccgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgtataga agcgtgtatg tatgaaagcg atttacttcg tggtgcattt 120  
cacttaggcg gctattacag gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagaac tccaaggcca gaaacagtac cagctccgag gtatggctac ctttgcgttaag 240  
tatcgtgagc agaaaagcggg atcgagtcta attaaacacg ctgaagaaat tttcgtaag 300  
agggggccgg acttgctttg gtgtaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gaaacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 238  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 238  
atgcttagagg taaaaccgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgtataga agcgtgtatg tatgaaagcg atttacttcg tggtgcattt 120  
cacttaggcg gctattacag gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cacccagaac tccaaggcca gaaacagtac cagctccgag gtatggctac ctttgcgttaag 240  
tatcgtgagc agaaaagcggg atcgagtcta attaaacacg ctgaagaaat tttcgtaag 300  
agggggccgg acttgctttg gtgtaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gaaacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 239  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 239  
atgcttagagg taaaaccgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgtataga agcgtgtatg tatgaaagcg atttacttcg tggtgcattt 120  
cacttaggcg gctattacgg gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac ctttgcgttaag 240  
tatcgtgagc agaaaagcggg atcgagtcta attaaacacg ctgaagaaat tttcgtaag 300  
agggggccgg acttgctttg gtgtaatgcg cggacatccg cctcaggcta ctacaaaaag 360

ttaggcttca gcgagcaggg agagatattt gaaacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 240  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 240  
atgcttagagg taaaaccat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agacccaaacc agccgataga agtgttatg tatgaaagcg atttacttcg tgggcattt 120  
cacttaggcg gctttacgg gggcaactg atttccatag cttcattcca ccaggccgag 180  
cactcagaac tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcggg atcgagtcta attaaacacg ctgaagaaat tcttcgttaag 300  
agggggcgg acttgctttg gtgtaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gaaacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 241  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 241  
atgatagaag taaaacctat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agacccaaacc agccgataga agtgttatg tatgaaacccg atttacttcg tgggcattt 120  
cacttaggcg gctttacgg gggcaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgatc agaaaagcggg atcgagtcta attagacacg ctgaacaaaat tcttcgttaag 300  
agggggcgg acatgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gaaacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 242  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 242  
atgatagaag taaaacctat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agacccaaacc agccgtttaga agtgttatg tatgaaacccg atttacttcg tgggcattt 120  
cacttaggcg gctttacgg gggcaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgatc agaaaagcggg atcgagtcta attagacacg ctgaacaaaat tcttcgttaag 300  
agggggcgg acatgctttg gtgtaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gaaacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 243  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<221> misc\_feature  
<222> 9, 76, 98  
<223> n = A,T,C or G

<400> 243  
atgcttagang taaaaccgat taacgcagag gatacctatg aactaaggca taaaataactc 60  
agaccaaacc agccgntaga agtgttatg tatgaaancg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcggtatc agaaaagcggg atcgagtcta attaaacacg ctgaacaaat tcttcgtgag 300  
agggggcgg acatgcttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gacacgcccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 244  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 244  
atgcttagaag taaaaccstat taacgcagag gatacctatg aactaaggca taaaataactc 60  
agaccaaacc agccgataga agtgttatg tatgaaacccg atttacttcg tgggcattt 120  
cacttaggcg gctttacgg gggcaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcggtatc agaaaagcggg atcgagtcta attagacacg ctgaacaaat tcttcgttaag 300  
agggggcgg acatgcttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gaaacgcccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 245  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 245  
atgcttagagg taaaaccgat taacgcagag gatacctatg aactaaggca taaaataactc 60  
agaccaaacc agccgttaga agtgttatg tatgaaacccg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaactg atttccatag cgtcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcggtatc agaaaagcggg atcgagtcta attagacacg ctgaacaaat tcttcgttaag 300  
agggggcgg acatgcttg gtgcaatgcg cggacatccg cctcaggcta ctacagaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gaaacgcccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 246  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 246  
atgatagagg taaaaccgat taacgcagag gatacctatg aactaaggca taaaatactc 60  
agaccaaacc agccgttaga agtgttatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgatc agaaaagcggg atcgagtcta attagacacg ctgaacaaat tttcgtaag 300  
agggggccgg acatgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg ggaggtattt gaaacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 247  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 247  
atgatagagg taaaaccgat taacgcagag gatacctatg aactaaggca taaaatactc 60  
agaccaaacc agccgttaga agtgttatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcggg atcgagtcta attagacacg ctgaacaaat tttcgtaag 300  
agggggccgg acatgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gaaacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 248  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 248  
atgatagagg taaaaccgat taacgcagag gatacctatg aactaaggca taaaatactc 60  
agaccaaacc agccgttaga agtgttatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcggg atcgagtcta attagacacg ctgaacaaat tttcgtaag 300  
agggggccgg acatgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gaaacgccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 249  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 249  
atgcttagagg taaaaccgat taacgcagag gatacctatg aactaaggca taaaatactc 60  
agaccaaacc agccgttaga agtgttatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctttacgg gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcggacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgatc agaaaagcggg atcgagtcta attagacacg ctgaagaaat tttcgtaag 300

aggggggcgg acatgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gacacgcccgc cagtaggacc tcacatcctg 420  
atgtataaaa agatcacata a 441

<210> 250  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 250  
atgctagaag taaaaccgat taacgcagag gatacctatg aactaaggca taaaataactc 60  
agaccaaacc agccgttaga agtgtgtatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctttacgg gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgatc agaaaagcggg atcgactcta attaaacacg ctgaacaaat tttcgtaag 300  
agggggcgg acatgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gaaacgcccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 251  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 251  
atgatagagg taaaaccgat taacgcagag gatacctatg aactaaggca tagaataactc 60  
agaccaaacc agccgttaga agtgtgtatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcggg atcgagtcta attaaacacg ctgaagaaat tttcgtaag 300  
agggggcgg acttgctttg gtgtaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agagatattt gatacggccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggatcacata a 441

<210> 252  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic DNA Sequence

<400> 252  
atgatagagg taaaaccgat taacgcagag gatacctatg aactaaggca tagagtactc 60  
agaccaaacc agccgttaga agtgtgtatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctattacag gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagaac tccaaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcggg atcgagtcta attaaacacg ctgaagaaat tttcgtaag 300  
agggggcgg acttgctttg gtgcaatgcg cggacatccg cctcaggcta ctacaaaaag 360  
ttaggcttca gcgagcaggg agaggtattt gagacggccgc cagtaggacc tcacatcctg 420  
atgtataaaa ggctcacgt a 441

<210> 253  
<211> 441  
<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 253

atgcttagagg taaaaccgat taacgcagag gataacttacg aactaaggca taaaatactc 60  
agaccaaacc agccgataga agtgttatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctattacag gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagaac tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcg 300  
agggggcg 360  
ttaggcttca gcgagcaggg agagatattt gatacgcgc ctaggacc tcacatcctg 420  
atgtataaaa ggatcacgta a 441

<210> 254

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 254

atgcttagagg taaaaccgat taacgcagag gataacctatg aactaaggca tagaatactc 60  
agaccaaacc agccgataga agtgttatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctttacag gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagaac tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcg 300  
agggggcg 360  
ttaggcttca gcgagcaggg agagatattt gaaacgcgc ctaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 255

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 255

atgcttagagg taaaaccgat taacgcagag gataacctatg aactaaggca taaaatactc 60  
agaccaaacc agccgttaga agtgttatg tatgaaaccg atttacttcg tgggcattt 120  
cacttaggcg gctattacag gggcaaactg atttccatag cttcattcca ccaggccgag 180  
cactcagacc tccaaggcca gaaacagtac cagctccgag gtatggctac cttggaaggt 240  
tatcgtgagc agaaaagcg 300  
agggggcg 360  
ttaggcttca gcgagcaggg agaggtattt gatacgcgc ctaggacc tcacatcctg 420  
atgtataaaa ggctcacata a 441

<210> 256

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 256

atgattgaag tcaaaccatat aaacgcggaa gatacgttatg agatcaggca ccgcattctc 60

cggccgaatc agccgctgga agcatgcaag tatgaaaccg atttgctcg gggcacgtt 120  
cacctcggtg gatattacg gggcaagctg atcagcatcg cttcccttca taatgccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaagga 240  
taccgtgagc aaaaagcggg aagcacgctc atccgccatg ccgaagagct tcttcggaaa 300  
aaaggcgcgg accttttatg gtgcaacgccc aggacatctg tgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgatcggacc tcataaaaa 420  
atgtataaga aattgacgta a 441

<210> 257

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 257

atgattgaag tcaaaccat aaacgcggaa gatacgtatg agatcaggca ccgcatttctc 60  
cggccgaatc agccgcttga agcatgtatg tatgaaaccg atttgctcg gggcacgtt 120  
cacctcggtg gatattacg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtcggc aaaaagcggg cagtagctt atccgccatg ccgaagagct tcttcggaaa 300  
aaggggccag accttttatg gtgcaatgcc aggacatctg tgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgatcggacc tcataaaaa 420  
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<210> 258

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 258

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cacctcggtg gatattacg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtcggc aaaaagcggg tagtacgctt atccgccatg ccgaagagct tcttcggaaa 300  
aaggggccag accttttatg gtgcaacgccc aggacatctg tgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaagg cggggctcgc gatataccgc cgatcggacc tcataaaaa 420  
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<210> 259

<211> 441

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 259

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cacctcggtg gatattacg gggcaagctg atcagcatcg cttcccttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtcggc aaaaagcggg cagtagctt atccgccatg ccgaagagct tcttcggaaa 300  
aaggggccag accttttatg gtgcaacgccc aggacatctg tgagcggcta ctataaaaag 360  
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atgtataaga aattgacata a 441

<210> 260  
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<212> DNA  
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<220>  
<223> Synthetic DNA Sequence

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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcctttca taatgccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac gcttgaaggg 240  
taccgcgagc aaaaagcggg aagcacgctc atccgccatg ccgaagagct tcttcggaaa 300  
aaaggcgcag accttttatg gtgcaacgccc aggacatctg tgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaggg cgaagtctac gacataccgc cgatcggacc tcataaaaa 420  
atgtataaga aattgacgta a 441

<210> 261  
<211> 441  
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<213> Artificial Sequence

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cgccgcgaatc agccgcttga agcatgtatg tatgaaaccg atttgctcgg gggcacgttt 120  
cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcctttca tcaagccgaa 180  
catccagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgcgagc aaaaagcggg cagtacgctt atccgccatg ccgaagagct tcttcggaaa 300  
aaaggcgcag accttttatg gtgcaacgccc aggacatctg tgagcggcta ctatgaaaag 360  
ctcggcttca gcbaacaggg cgaagtctgc gacataccgc cgatcggacc tcataaaaa 420  
atgtataaga aattgacata a 441

<210> 262  
<211> 441  
<212> DNA  
<213> Artificial Sequence

<220>  
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<400> 262  
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cacctcggtg gatattaccg gggcaagctg atcagcatcg cttcctttca tcaagccgaa 180  
cattcagagc ttgaaggcca aaaacagtat cagctgagag ggatggcgac acttgaaggg 240  
taccgtgagc aaaaagcggg cagtacgctt atccgccatg ccgaagagct tcttcggaaa 300  
aagggggcag accttttatg gtgcaacgccc aggacatctg tgagcgggta ctataaaaag 360  
ctcggcttca gcbaacaagg cggggctac gatataccgc cgatcggacc tcataaaaa 420  
atgtataaga aattgacgta a 441

<210> 263  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 263

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 264

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 264

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Arg Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 265

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 265

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Ser Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe Asn Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 266

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 266

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 267

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 267

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Ala  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Ala Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 268

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 268

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 269

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 269

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Leu His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 270

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 270

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 271

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 271

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
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His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Leu	Glu	Ala	Cys	Lys	Tyr	Glu
				20					25				30		
Thr	Asp	Leu	Leu	Gly	Gly	Thr	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
				35					40			45			
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Pro	Glu	Leu
				50					55		60				
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
				65				70		75			80		
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
				85					90			95			
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr	
				100				105				110			
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
				115				120			125				
Val	Tyr	Asp	Thr	Pro	Pro	Val	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
				130				135			140				
Leu	Thr														
	145														

<210> 272

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 272

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
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His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Leu	Glu	Ala	Cys	Lys	Tyr	Glu
				20					25			30			
Thr	Asp	Leu	Leu	Gly	Gly	Thr	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
				35					40		45				
Lys	Leu	Val	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Ser	Glu	Leu
				50					55		60				
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Glu
				65				70		75			80		
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
				85					90			95			
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr	
				100				105			110				
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
				115				120			125				
Val	Tyr	Asp	Ile	Pro	Pro	Val	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
				130				135			140				
Leu	Thr														
	145														

<210> 273

<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 273

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 274  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 274

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Arg Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 275  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 275  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Leu His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 276  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 276  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 277  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

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<400> 277
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg
   1           5           10          15
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu
   20          25          30
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly
   35          40          45
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu
   50          55          60
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu
   65          70          75          80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu
   85          90          95
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr
   100         105         110
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu
   115         120         125
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys
   130         135         140
Leu Thr
   145

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<210> 278  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

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<400> 278
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg
   1           5           10          15
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu
   20          25          30
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly
   35          40          45
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu
   50          55          60
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly
   65          70          75          80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu
   85          90          95
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr
  100         105         110
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu
  115         120         125
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys
  130         135         140
Leu Thr
  145

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<210> 279  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 279

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 280  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 280

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr

<210> 281  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 281  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Leu His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 282  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 282  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140

Leu Thr  
145

<210> 283  
<211> 146  
<212> PRT  
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<220>  
<223> Synthetic Protein Sequence

<400> 283  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Gln Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 284  
<211> 146  
<212> PRT  
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<220>  
<223> Synthetic Protein Sequence

<400> 284  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Gln Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys

	130	135	140
Leu	Ala		
	145		

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<212> PRT  
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<220>  
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<400> 285

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His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Leu	Glu	Ala	Cys	Met	Tyr	Glu
				20				25				30			
Thr	Asp	Leu	Leu	Gly	Gly	Ala	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Gln	Gly
				35			40			45					
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Lys	Ala	Glu	His	Ser	Glu	Leu
				50			55			60					
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
				65			70		75			80			
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
				85			90			95					
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr	
				100			105				110				
Ser	Val	Arg	Gly	Tyr	Tyr	Glu	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Gly
				115			120			125					
Val	Tyr	Asp	Ile	Pro	Pro	Ile	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
				130			135			140					
Leu	Thr														
	145														

<210> 286  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
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<400> 286

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
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His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Leu	Glu	Ala	Cys	Lys	Tyr	Glu
					20			25				30			
Thr	Asp	Leu	Leu	Gly	Gly	Thr	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
				35			40			45					
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Pro	Glu	Leu
				50			55			60					
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
				65			70		75			80			
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
				85			90			95					
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr	
				100			105				110				
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
				115			120			125					

Val His Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 287  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 287  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 288  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 288  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu

115                   120                   125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130                   135                   140  
Leu Thr  
145

<210> 289  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 289  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20                   25                   30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35                   40                   45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50                   55                   60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65                   70                   75                   80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85                   90                   95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100                  105                  110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115                  120                  125  
Val Tyr Asp Ala Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130                  135                  140  
Leu Thr  
145

<210> 290  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 290  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20                   25                   30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35                   40                   45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50                   55                   60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65                   70                   75                   80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85                   90                   95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100                  105                  110

Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 291  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 291  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

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<211> 146  
<212> PRT  
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<220>  
<223> Synthetic Protein Sequence

<400> 292  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr

100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 293  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 293  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Ala  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 294  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 294  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95

Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 295  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
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<400> 295  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 296  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 296  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Gln Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu

	85		90		95										
Leu	Leu	Arg	Lys	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr
			100			105						110			
Ser	Val	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Gly
			115			120						125			
Ile	Tyr	Asp	Ile	Pro	Pro	Ile	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
			130			135						140			
Leu	Thr														
	145														

<210> 297

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 297

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
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His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Leu	Glu	Ala	Cys	Lys	Tyr	Glu
								20		25			30		
Thr	Asp	Leu	Leu	Gly	Gly	Thr	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Gln	Gly
						35		40			45				
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Lys	Ala	Glu	His	Ser	Glu	Leu
						50		55			60				
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
					65		70		75			80			
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
						85		90			95				
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr	
						100		105			110				
Ser	Val	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
						115		120			125				
Val	Tyr	Asp	Ile	Pro	Pro	Ile	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
						130		135			140				
Leu	Thr														
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<210> 298

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 298

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
1			5			10			15					15	
His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Leu	Glu	Ala	Cys	Met	Tyr	Glu
						20		25			30				
Thr	Asp	Leu	Leu	Gly	Gly	Ala	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
						35		40			45				
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Ser	Glu	Leu
						50		55			60				
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
					65		70		75			80			

Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 299  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 299  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe Asn Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met His Lys Lys  
130 135 140  
Leu Thr  
145

<210> 300  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 300  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Ser Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe Asn Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly

65                   70                   75                   80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
      85              90              95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
      100            105            110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
      115            120            125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
      130            135            140  
Leu Thr  
145

<210> 301  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 301  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
  20            25            30  
Thr Asp Leu Leu Arg Ser Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
  35            40            45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
  50            55            60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
  65            70            75            80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
  85            90            95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
  100           105           110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
  115           120           125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
  130           135           140  
Leu Thr  
145

<210> 302  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 302  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
  20            25            30  
Thr Asp Leu Leu Gly Ser Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
  35            40            45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
  50            55            60

Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 303

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 303

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Arg Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 304

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 304

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Ser Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe Asn Gln Ala Glu His Pro Glu Leu

50               55               60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65               70               75               80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85               90               95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100              105              110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115              120              125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130              135              140  
Leu Thr  
145

<210> 305  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 305  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1               5               10               15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20              25              30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35              40              45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50              55              60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65              70              75              80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85              90              95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100             105             110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115             120             125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130             135             140  
Leu Thr  
145

<210> 306  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 306  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1               5               10               15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20              25              30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35              40              45

Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 307  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 307  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 308  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 308  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly

35	40	45
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu		
50	55	60
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly		
65	70	75
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu		80
85	90	95
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr		
100	105	110
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu		
115	120	125
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys		
130	135	140
Leu Thr		
145		

<210> 309

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 309

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg		
1	5	10
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu		15
20	25	30
Thr Asp Leu Leu Gly Ser Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly		
35	40	45
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu		
50	55	60
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu		
65	70	75
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Ala		80
85	90	95
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr		
100	105	110
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu		
115	120	125
Val Tyr Asp Thr Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys		
130	135	140
Leu Thr		
145		

<210> 310

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 310

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg		
1	5	10
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu		15
20	25	30

Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Ala  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Arg Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 311  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 311  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 312  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 312  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu

	20	25	30												
Thr	Asp	Leu	Leu	Gly	Gly	Thr	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
			35		40							45			
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	Asn	Gln	Ala	Glu	His	Pro	Glu	Leu
			50		55						60				
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
			65		70				75			80			
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
			85		90							95			
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr	
														100	
														105	
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Gly
														115	
														120	
Val	Tyr	Asp	Ile	Pro	Pro	Val	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
														130	
														135	
Leu	Thr													140	
															145

<210> 313

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 313

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
						1	5		10				15		
His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Leu	Glu	Ala	Cys	Lys	Tyr	Glu
									20	25			30		
Thr	Asp	Leu	Leu	Arg	Gly	Ala	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
								35	40			45			
Lys	Leu	Val	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Pro	Glu	Leu
									50	55		60			
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
								65	70		75		80		
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
								85	90			95			
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr	
									100	105		110			
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
								115	120			125			
Val	Tyr	Asp	Ile	Pro	Pro	Thr	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
								130		135			140		
Leu	Thr														145

<210> 314

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 314

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
						1	5		10			15			

His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Ala  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Ala Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 315  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 315  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Ser Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 316  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 316  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg

1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe Asn Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Val Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 317

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 317

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Met Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 318

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 318

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe Asn Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Ala  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 319

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 319

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Leu Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 320

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 320  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Ala Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 321  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 321  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Lys His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 322  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 322  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe Asn Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Ala  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 323  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 323  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 324  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 324  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Ala  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 325

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 325  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Thr Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 326

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 326

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 327

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 327

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe Asn Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 328

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 328

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Ala  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 329

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 329

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Ala  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 330

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 330

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 331

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 331

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Lys  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 332

<211> 146

<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 332  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 333  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 333  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Val Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Ala Gly Pro His Ile Leu Met Tyr Thr Lys  
130 135 140  
Leu Thr  
145

<210> 334

<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 334  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 335  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 335  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Lys Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe Asn Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Ala  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 336  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 336  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 337  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 337  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 338  
<211> 146  
<212> PRT  
<213> Artificial Sequence  
  
<220>  
<223> Synthetic Protein Sequence  
  
<400> 338  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Lys  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 339  
<211> 146  
<212> PRT  
<213> Artificial Sequence  
  
<220>  
<223> Synthetic Protein Sequence  
  
<400> 339  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 340  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 340  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Glu Glu Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg Tyr Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 341  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 341  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Leu His Leu Gly Gly Tyr Tyr Gln Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Gly Leu  
50 55 60  
Glu Gly Glu Glu Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr

<210> 342  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 342  
Met Ile Asp Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Gln Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 343  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 343  
Met Ile Glu Val Lys Pro Ile Ser Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Ala Phe His Leu Gly Gly Tyr Tyr Gln Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140

Leu Thr  
145

<210> 344  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 344  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Ala  
145

<210> 345  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 345  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Cys Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Arg Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys

130  
Leu Ala  
145

135

140

<210> 346  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 346  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Cys Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Gln Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 347  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 347  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Gln Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Asn Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Val Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125

Ile Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Ala  
145

<210> 348  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 348  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 349  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 349  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Ala Phe His Leu Gly Gly Tyr Tyr Gln Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu

115                    120                    125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130                    135                    140  
Leu Thr  
145

<210> 350  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 350  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1                    5                    10                    15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20                    25                    30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Gln Gly  
35                    40                    45  
Lys Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Glu Leu  
50                    55                    60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65                    70                    75                    80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85                    90                    95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100                    105                    110  
Ser Ala Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115                    120                    125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130                    135                    140  
Leu Ala  
145

<210> 351  
<211> 146  
<212> PRT  
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<220>  
<223> Synthetic Protein Sequence

<400> 351  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20                    25                    30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Asp  
35                    40                    45  
Arg Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50                    55                    60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65                    70                    75                    80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85                    90                    95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100                    105                    110

Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 352  
<211> 146  
<212> PRT  
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<220>  
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<400> 352  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Ile  
100 105 110  
Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Ala  
145

<210> 353  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 353  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Asn Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr

	100	105	110
Ser Ala Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu			
115	120	125	
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys			
130	135	140	
Leu Ala			
145			

<210> 354  
<211> 146  
<212> PRT  
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<220>  
<223> Synthetic Protein Sequence

<400> 354  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Asp  
35 40 45  
Arg Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 355  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 355  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Gln Gly  
35 40 45  
Lys Leu Ile Ser Thr Ala Ser Phe His Gln Ala Gly His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Arg Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95

Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Ile  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 356  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 356  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Phe Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Ala Phe His Leu Gly Gly Tyr Tyr Gln Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 357  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 357  
Met Ile Glu Val Arg Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu

	85		90		95										
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr	
			100				105					110			
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
			115				120					125			
Ala	Tyr	Asp	Ile	Pro	Pro	Ile	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
			130				135					140			
Leu	Thr														
	145														

<210> 358

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 358

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
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His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Leu	Glu	Ala	Cys	Met	Tyr	Glu
					20				25				30		
Thr	Asp	Leu	Leu	Gly	Gly	Thr	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
					35			40				45			
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Ser	Glu	Leu
					50			55			60				
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
					65			70			75		80		
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
					85			90			95				
Leu	Leu	Arg	Lys	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr
					100			105				110			
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Gly
					115			120				125			
Val	Tyr	Asp	Ile	Pro	Pro	Ile	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
					130			135				140			
Leu	Thr														
	145														

<210> 359

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 359

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
1					5				10					15	
His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Leu	Glu	Ala	Cys	Met	Tyr	Glu
					20				25			30			
Thr	Asp	Leu	Leu	Gly	Gly	Thr	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
					35			40			45				
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Ser	Glu	Leu
					50			55			60				
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
					65			70			75		80		

Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 360  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 360  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Pro Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Ile  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 361  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 361  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly

65                   70                   75                   80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
              85               90               95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
              100              105              110  
Ser Ala Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Lys Gln Gly Glu  
              115              120              125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
              130              135              140  
Leu Thr  
145

<210> 362  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 362  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1               5               10               15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20              25              30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35              40              45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50              55              60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65              70              75              80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85              90              95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Ile  
100             105             110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115             120             125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130             135             140  
Leu Thr  
145

<210> 363  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 363  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1               5               10               15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20              25              30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35              40              45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50              55              60

Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Gly Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 364

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 364

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Gln Asp  
35 40 45  
Arg Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg Tyr Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Ile  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 365

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 365

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu

50               55               60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65               70               75               80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85               90               95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100              105              110  
Ser Ala Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115              120              125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130              135              140  
Leu Thr  
145

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<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 366  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20              25              30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35              40              45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50              55              60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65              70              75              80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85              90              95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100             105             110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115             120             125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130             135             140  
Leu Thr  
145

<210> 367  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 367  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20              25              30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35              40              45

Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Ile  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 368

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 368

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro Tyr Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 369

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 369

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly

35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 370  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 370  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 371  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 371  
Met Ile Glu Val Asn Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30

Thr Asp Leu Leu Gly Gly Thr Ser His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Asn Ala Glu His Ser Glu Leu  
50 55 60  
Asp Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Ser Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 372  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 372  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Ala  
145

<210> 373  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 373  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu

	20	25	30												
Thr	Asp	Leu	Leu	Gly	Gly	Ala	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
			35		40							45			
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ser	Glu	His	Pro	Glu	Leu
			50		55						60				
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
			65		70				75				80		
Tyr	Arg	Glu	Leu	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
													85		95
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Ile	
													100		110
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
													115		125
Val	Tyr	Asp	Ile	Pro	Pro	Ile	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
												130		140	
Leu	Thr														
		145													

<210> 374

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 374

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
						5			10				15		
His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Leu	Glu	Ala	Cys	Met	Tyr	Glu
									20		25		30		
Thr	Asp	Leu	Leu	Gly	Gly	Thr	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
									35		40		45		
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Ser	Glu	Leu
									50		55		60		
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Glu
									65		70		75		80
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
									85		90		95		
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Ile	
									100		105		110		
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Gly
									115		120		125		
Val	Tyr	Asp	Ile	Pro	Pro	Ile	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
									130		135		140		
Leu	Thr														
		145													

<210> 375

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 375

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
						1			5				10		15

His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Gly Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 376  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 376  
Met Ile Glu Ala Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 377  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 377  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg

1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu His Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 378

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 378

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 379

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 379

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 380

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 380

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Ala Phe His Leu Gly Gly Tyr Tyr Gln Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Val His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 381

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 381  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly His Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 382  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 382  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 383  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 383  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 384  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 384  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Val His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 385  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 385  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Ala Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Asn Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Val Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 386

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 386  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Arg Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Ile  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 387

<211> 146

<212> PRT

<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<221> unsure  
<222> 17  
<223> Xaa = His or Pro

<400> 387

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Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg
   1          5           10          15
Xaa Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu
   20         25           30
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly
   35         40           45
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu
   50         55           60
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly
   65         70           75           80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu
   85         .            90           95
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr
  100        105          110
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly
  115        120          125
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys
  130        135          140
Leu Thr
  145

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<210> 388

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 388

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Met Ile Glu Ala Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg
   1          5          10          15
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu
   20         25          30
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly
   35         40          45
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu
   50         55          60
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly
   65         70          75          80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu
   85         90          95
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr
  100        105         110
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu
  115        120         125
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys
  130        135         140
Leu Thr
  145

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<210> 389  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 389

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145.

<210> 390  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 390

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 391  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 391  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Leu His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg Gln Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Glu  
130 135 140  
Leu Thr  
145

<210> 392  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 392  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 393  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 393

Met Ile Glu Val Lys Pro Ile Asn Ala Gly Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 394  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 394

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr

<210> 395  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 395  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Val Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Val Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 396  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 396  
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His Arg Ile Leu Arg Pro Asn Arg Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Gly Leu  
50 55 60  
Glu Gly Lys Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140

Leu Thr  
145

<210> 397  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Glu Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 398  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 398  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
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Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Asp Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys

130  
Leu Thr  
145

135

140

<210> 399  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Gly
   20         25           30
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly
   35         40           45
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu Gln Pro Glu Leu
   50         55           60
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly
   65         70           75           80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu
   85         90           95
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr
  100        105          110
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly
  115        120          125
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys
  130        135          140
Leu Thr
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<210> 400  
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				20					25					30	
Thr	Asp	Leu	Leu	Gly	Gly	Thr	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
						35			40					45	
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Ser	Glu	Leu
						50			55					60	
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
					65		70			75					80
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
							85			90					95
Ile	Leu	Arg	Lys	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr
							100			105					110
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Gly
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130 135 140  
Leu Thr  
145

<210> 401  
<211> 146  
<212> PRT  
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<220>  
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20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Pro Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 402  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu

115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 403  
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<212> PRT  
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<220>  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 404  
<211> 146  
<212> PRT  
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<220>  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110

Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

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<211> 146  
<212> PRT  
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<220>  
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20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 406  
<211> 146  
<212> PRT  
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<220>  
<223> Synthetic Protein Sequence

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20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Gly Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr

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Ser	Ala	Ser	Gly	Tyr	Tyr	Glu	Lys	Leu	Gly	Phe	Ser	Gly	Gln	Gly	Glu
	115		120			125									
Val	Tyr	Asp	Ile	Pro	Pro	Ile	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
	130		135								140				
Leu	Thr														
	145														

<210> 407  
<211> 146  
<212> PRT  
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<220>  
<223> Synthetic Protein Sequence

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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 408  
<211> 146  
<212> PRT  
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<220>  
<223> Synthetic Protein Sequence

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Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95

Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val His Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 409  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Ile  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 410  
<211> 146  
<212> PRT  
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<220>  
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20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu

	85		90		95										
Leu	Leu	Arg	Lys	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr
			100			105						110			
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Gly
			115			120						125			
Val	Tyr	Asp	Ile	Pro	Pro	Val	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
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Leu	Thr														
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<210> 411  
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<213> Artificial Sequence

<220>  
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20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
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<210> 412  
<211> 146  
<212> PRT  
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<220>  
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20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80

Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
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<210> 413  
<211> 146  
<212> PRT  
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<220>  
<223> Synthetic Protein Sequence

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20 25 30  
Thr Asp Leu Leu Gly Gly Thr Leu His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
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<210> 414  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

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20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Cys Ile Ala Ser Phe His Lys Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Asp Gly

65                   70                   75                   80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
              85               90               95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
              100           105           110  
Ser Val Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
              115           120           125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
              130           135           140  
Leu Thr  
145

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<211> 146  
<212> PRT  
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<220>  
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20              25              30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35              40              45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50              55              60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65              70              75              80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85              90              95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100           105           110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115           120           125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130           135           140  
Leu Thr  
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<211> 146  
<212> PRT  
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<220>  
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<400> 416  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
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20              25              30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35              40              45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50              55              60

Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 417  
<211> 146  
<212> PRT  
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<220>  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
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Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 418  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 418  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu

50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 419  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 419  
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His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Val Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 420  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 420  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45

Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 421  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 421  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 422  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 422  
Met Ile Glu Ala Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly

35                   40                   45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Thr Glu Leu  
50                   55                   60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65                   70                   75                   80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85                   90                   95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100                 105                 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115                 120                 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130                 135                 140  
Leu Thr  
145

<210> 423

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 423

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1                   5                   10                   15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Asp Ala Cys Lys Tyr Glu  
20                 25                 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35                 40                 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50                 55                 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65                 70                 75                   80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85                 90                 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100                105                110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115                120                125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130                135                140  
Leu Thr  
145

<210> 424

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 424

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1                   5                   10                   15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20                 25                 30

Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Ala  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 425

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 425

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Met Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr His Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asn Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 426

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 426

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu

	20	25	30												
Thr	Asp	Leu	Leu	Gly	Gly	Thr	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
		35			40							45			
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Pro	Glu	Leu
		50			55						60				
Val	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
		65			70				75				80		
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
														95	
					85				90						
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr	
													100		
									105				110		
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
													115		
									120				125		
Val	Tyr	Asp	Ile	Pro	Pro	Val	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
												130			
Leu	Thr										135			140	
		145													

<210> 427

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 427

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
						1	5		10				15		
His	Arg	Val	Leu	Arg	Pro	Asn	Gln	Pro	Leu	Glu	Ala	Cys	Met	Tyr	Glu
									20	25			30		
Thr	Asp	Leu	Leu	Gly	Gly	Thr	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
									35	40			45		
Glu	Leu	Val	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Pro	Glu	Leu
									50	55			60		
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
									65	70			75		80
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
									85	90			95		
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr	
									100	105			110		
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Gly
									115	120			125		
Val	Tyr	Asp	Ile	Pro	Pro	Val	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
									130				135		
Leu	Thr														
		145													

<210> 428

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 428

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
						1	5		10			15			

His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Asp Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 429  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 429  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Leu Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 430  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 430  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg

1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Leu His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Val Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Gly Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 431

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 431

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Ala  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 432

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 432

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Leu His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Ile Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 433

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 433

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe Asn Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 434

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 434  
Met Ile Glu Val Lys Pro Ile Thr Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Thr Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 435  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 435  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Ala Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Ser Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Glu  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 436  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 436

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg			
1	5	10	15
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu			
20	25	30	
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Gln Gly			
35	40	45	
Lys Leu Ile Ser Ile Ala Ser Phe His Asn Ala Glu His Ser Glu Leu			
50	55	60	
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly			
65	70	75	80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu			
85	90	95	
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr			
100	105	110	
Ser Val Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu			
115	120	125	
Val Tyr Asp Ile Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys			
130	135	140	
Leu Ala			
145			

<210> 437  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 437

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg			
1	5	10	15
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu			
20	25	30	
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly			
35	40	45	
Lys Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Glu Leu			
50	55	60	
Glu Gly Glu Glu Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly			
65	70	75	80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu			
85	90	95	
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr			
100	105	110	
Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu			
115	120	125	
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys			
130	135	140	
Leu Thr			
145			

<210> 438  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 438

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Thr Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Thr Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 439

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 439

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Gly Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Thr Leu Ile Lys His Ala Glu Gln  
85 90 95  
Leu Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Ile Thr  
145

<210> 440

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 440

Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Gln  
85 90 95  
Leu Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 441

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 441

Met Leu Glu Val Lys Leu Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Phe Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asn Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 442

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 442

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 443

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 443

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Gly Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Ile Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 444

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 444

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Gln  
85 90 95  
Leu Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 445

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 445

Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 446

<211> 146

<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 446  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Glu Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85 90 95  
Leu Leu Arg Lys Gly Ala Asn Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Thr Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 447  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 447  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Leu His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Phe Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Lys  
115 120 125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 448

<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 448

Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 449  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 449

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Phe Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 450  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 450  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Phe Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Gln  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 451  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 451  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Gly Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Lys Gly Ala Asn Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 452  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<221> unsure  
<222> 45  
<223> Xaa = Phe, Ser, Tyr, or Cys

<221> unsure  
<222> 105  
<223> Xaa = Leu, Met, or Val

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<400> 452
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg
      1           5           10          15
His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu
      20          25          30
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Xaa Tyr Arg Gly
      35          40          45
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu
      50          55          60
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly
      65          70          75          80
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Gln
      85          90          95
Ile Leu Arg Lys Arg Gly Ala Asp Xaa Leu Trp Cys Asn Ala Arg Thr
      100         105         110
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu
      115         120         125
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg
      130         135         140
Leu Thr
      145

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<210> 453  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

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<400> 453
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg
   1           5           10          15
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Val Cys Met Tyr Glu
   20          25          30
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly
   35          40          45
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu
   50          55          60
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly
   65          70          75          80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Gln
   85          90          95

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Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 454  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 454  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Phe Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 455  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 455  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Val Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Phe Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln

	85	90	95												
Ile	Leu	Arg	Lys	Arg	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr
			100					105						110	
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
			115					120					125		
Ile	Phe	Asp	Thr	Pro	Pro	Val	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Arg
	130					135						140			
Leu	Thr														
145															

<210> 456  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 456

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Leu	Arg
1				5				10					15		
His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Ile	Glu	Ala	Cys	Met	Phe	Glu
					20			25					30		
Ser	Asp	Leu	Leu	Arg	Gly	Ala	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
					35			40				45			
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Ser	Asp	Leu
					50			55			60				
Gln	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
					65			70			75		80		
Tyr	Arg	Asp	Gln	Lys	Ala	Gly	Ser	Ser	Leu	Ile	Arg	His	Ala	Glu	Gln
					85			90			95				
Ile	Leu	Arg	Lys	Arg	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr
					100			105				110			
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
					115			120				125			
Ile	Phe	Asp	Thr	Pro	Pro	Val	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Arg
	130					135						140			
Leu	Thr														
145															

<210> 457  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 457

Met	Leu	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Leu	Arg
1				5				10					15		
His	Lys	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Leu	Glu	Val	Cys	Met	Tyr	Glu
					20			25				30			
Thr	Asp	Leu	Leu	Arg	Gly	Ala	Phe	His	Leu	Gly	Gly	Phe	Tyr	Arg	Gly
					35			40			45				
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Ser	Glu	Leu
					50			55			60				
Gln	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
					65			70			75		80		

Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 458  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 458  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Ile Glu Val Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Gln  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 459  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 459  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Phe Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly

65                   70                   75                   80  
Phe Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
              85               90               95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
              100              105              110  
Ser Ala Ser Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu Gln Gly Glu  
              115              120              125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
              130              135              140  
Leu Thr  
145

<210> 460  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 460  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1               5               10               15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Phe Glu  
20              25              30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35              40              45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50              55              60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65              70              75              80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85              90              95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100             105             110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115             120             125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130             135             140  
Ile Thr  
145

<210> 461  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 461  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1               5               10               15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu  
20              25              30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Gly Gly  
35              40              45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Asp Leu  
50              55              60

Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85 90 95  
Leu Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 462  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 462  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Leu His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 463  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 463  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Phe Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Gly Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu

50               55               60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65               70               75               80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85               90               95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100              105              110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115              120              125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130              135              140  
Ile Thr  
145

<210> 464  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 464  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1               5               10               15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Val Cys Met Tyr Glu  
20              25              30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35              40              45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50              55              60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65              70              75              80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Gln  
85              90              95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100             105             110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115             120             125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130             135             140  
Leu Thr  
145

<210> 465  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 465  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1               5               10               15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Val Cys Met Tyr Glu  
20              25              30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Gly Gly  
35              40              45

Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 466  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 466  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Glu Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Gln Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Phe Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Glu  
85 90 95  
Ile Leu Arg Asn Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Thr Ala Ser Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu His Gly Glu  
115 120 125  
Val Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 467  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 467  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly

35                   40                   45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50                   55                   60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65                   70                   75                   80  
Phe Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Glu  
85                   90                   95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100                105                110  
Thr Ala Ala Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115                120                125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130                135                140  
Ile Thr  
145

<210> 468

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 468

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1                   5                   10                   15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20                25                30  
Ser Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly  
35                40                45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50                55                60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65                70                75                   80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85                90                95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100               105               110  
Thr Ala Ala Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu Gln Gly Glu  
115               120               125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130               135               140  
Leu Thr  
145

<210> 469

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 469

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1                   5                   10                   15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20                25                30

Ser Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu Gln Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Thr Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ala Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 470

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 470

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Glu Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Phe Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 471

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 471

Met Leu Glu Val Lys Pro Ile Asn Ala Glu Glu Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu

	20	25	30
Thr Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly			
35	40	45	
Gln Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu			
50	55	60	
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly			
65	70	75	80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Lys His Ala Glu Glu			
85	90	95	
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr			
100	105	110	
Thr Ala Ala Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu			
115	120	125	
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys			
130	135	140	
Ile Thr			
145			

<210> 472

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 472

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg			
1	5	10	15
His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu			
20	25	30	
Ser Asp Leu Leu Arg Ser Ala Phe His Leu Gly Gly Phe Tyr Arg Gly			
35	40	45	
Lys Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Glu Leu			
50	55	60	
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly			
65	70	75	80
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Glu			
85	90	95	
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr			
100	105	110	
Ser Ala Ala Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu Gln Gly Glu			
115	120	125	
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg			
130	135	140	
Ile Thr			
145			

<210> 473

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 473

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg			
1	5	10	15

His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Gln  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ala Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu His Gly Glu  
115 120 125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 474

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 474

Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Thr Ala Ala Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Tyr Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 475

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 475

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Glu Thr Tyr Glu Leu Arg

1	5	10	15
His Lys Ile Leu Arg Pro Asn Gln Pro	Leu Glu Ala Cys Met Tyr Glu		
20	25	30	
Thr Asp Leu Leu Arg Gly Ala Phe His	Leu Gly Gly Phe Tyr Arg Gly		
35	40	45	
Lys Leu Ile Ser Ile Ala Ser Phe His	Gln Ala Asp His Ser Glu Leu		
50	55	60	
Gln Gly Gln Lys Gln Tyr Gln Leu Arg	Gly Met Ala Thr Leu Glu Gly		
65	70	75	80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr	Leu Ile Arg His Ala Glu Gln		
85	90	95	
Ile Leu Arg Lys Arg Gly Ala Asp	Leu Leu Trp Cys Asn Ala Arg Thr		
100	105	110	
Ser Ala Ala Gly Tyr Tyr Lys Lys	Leu Gly Phe Ser Glu His Gly Glu		
115	120	125	
Ile Phe Glu Thr Pro Pro Val	Gly Pro His Ile Leu Met Tyr Lys Arg		
130	135	140	
Leu Thr			
145			

<210> 476

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 476

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg			
1	5	10	15
His Arg Ile Leu Arg Pro Asn Gln Pro	Ile Glu Ala Cys Met Tyr Glu		
20	25	30	
Ser Asp Leu Leu Arg Gly Ala Phe His	Leu Gly Gly Phe Tyr Arg Gly		
35	40	45	
Lys Leu Ile Ser Ile Ala Ser Phe His	Gln Ala Glu His Ser Asp Leu		
50	55	60	
Gln Gly Gln Lys Gln Tyr Gln Leu Arg	Gly Met Ala Thr Leu Glu Gly		
65	70	75	80
Tyr Arg Glu Gln Lys Ala Gly Ser Ser	Leu Ile Arg His Ala Glu Glu		
85	90	95	
Ile Leu Arg Lys Lys Gly Ala Asp	Met Leu Trp Cys Asn Ala Arg Thr		
100	105	110	
Thr Ala Ala Gly Tyr Tyr Lys Arg	Leu Gly Phe Ser Glu Gln Gly Glu		
115	120	125	
Val Phe Asp Thr Pro Pro Val	Gly Pro His Ile Leu Met Tyr Lys Arg		
130	135	140	
Ile Thr			
145			

<210> 477

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 477

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Gln Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Gln Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Phe Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ala Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu His Gly Glu  
115 120 125  
Ile Phe Asp Thr Pro Pro Ala Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 478

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 478

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Glu Thr Tyr Glu Leu Arg  
1 5 10 15  
Gln Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Gln Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ala Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu His Gly Glu  
115 120 125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 479

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 479  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Gln Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Lys His Ala Glu Gln  
85 90 95  
Leu Leu Arg Glu Lys Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ala Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 480  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 480  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Asp Leu  
50 55 60  
Glu Gly Gln Asn Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu His Gly Glu  
115 120 125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 481  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 481  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Gln Leu Ile Ser Ile Ala Ser Phe His Lys Ala Glu His Ser Asp Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Gln  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ala Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 482  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 482  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Gln  
85 90 95  
Leu Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu His Gly Glu  
115 120 125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 483  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 483

Met Leu Glu Val Lys Pro Ile Asn Ala Glu Glu Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Gln Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Phe Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Lys His Ala Glu Gln  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ala Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu His Gly Glu  
115 120 125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 484

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 484

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Glu Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Thr Leu Ile Lys His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ala Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu His Gly Glu  
115 120 125  
Ile Tyr Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Ile Thr  
145

<210> 485

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 485

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ser Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Gln Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ala Gly Tyr Tyr Lys Arg Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Ile Thr  
145

<210> 486

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 486

Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Val Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 487

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 487

Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 488

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 488

Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Gly Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 489

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 489

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Leu	Arg
1				5					10					15	
His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Ile	Glu	Ala	Cys	Met	Tyr	Glu
				20				25					30		
Ser	Asp	Leu	Leu	Arg	Gly	Ala	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
				35			40			45					
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Ser	Glu	Leu
				50			55			60					
Gln	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
				65			70		75			80			
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Ser	Leu	Ile	Lys	His	Ala	Glu	Glu
				85			90			95					
Ile	Leu	Arg	Lys	Arg	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr
				100			105					110			
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
				115			120				125				
Ile	Phe	Glu	Thr	Pro	Pro	Val	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Arg
				130			135			140					
Ile	Thr														
	145														

<210> 490

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 490

Met	Leu	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Leu	Arg
1				5					10					15	
His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Ile	Glu	Ala	Cys	Met	Tyr	Glu
				20				25					30		
Ser	Asp	Leu	Leu	Arg	Gly	Ala	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
				35			40		45						
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Pro	Glu	Leu
				50			55		60						
Gln	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
				65			70		75			80			
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Ser	Leu	Ile	Lys	His	Ala	Glu	Glu
				85			90			95					
Ile	Leu	Arg	Lys	Arg	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr
				100			105				110				
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
				115			120			125					
Val	Phe	Glu	Thr	Pro	Pro	Val	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Arg
				130			135			140					
Ile	Thr														
	145														

<210> 491

<211> 146

<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 491  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Ala Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Gly Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 492  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 492  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Val Cys Met Tyr Glu  
20 25 30  
Ser Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 493

<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 493

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Val Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Gly Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 494  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 494

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Gly Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 495  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<221> unsure  
<222> 3  
<223> Xaa = Asp or Glu

<221> unsure  
<222> 26  
<223> Xaa = Leu, Ile, or Val

<221> unsure  
<222> 33  
<223> Xaa = Ile, Thr, Asn, or Ser

<400> 495

Met	Leu	Xaa	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Leu	Arg
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His	Lys	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Xaa	Glu	Val	Cys	Met	Tyr	Glu
				20				25				30			
Xaa	Asp	Leu	Leu	Arg	Gly	Ala	Phe	His	Leu	Gly	Gly	Phe	Tyr	Arg	Gly
				35			40			45					
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Ser	Asp	Leu
				50		55			60						
Gln	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
				65		70		75		80					
Tyr	Arg	Asp	Gln	Lys	Ala	Gly	Ser	Ser	Leu	Ile	Lys	His	Ala	Glu	Gln
				85		90			95						
Ile	Leu	Arg	Glu	Arg	Gly	Ala	Asp	Met	Leu	Trp	Cys	Asn	Ala	Arg	Thr
				100		105			110						
Ser	Ala	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
				115		120		125							
Val	Phe	Asp	Thr	Pro	Pro	Val	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Arg
				130		135			140						
Leu	Thr														
	145														

<210> 496  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 496

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1				5				10				15			
His	Lys	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Ile	Glu	Val	Cys	Met	Tyr	Glu
				20				25			30				
Thr	Asp	Leu	Leu	Arg	Gly	Ala	Phe	His	Leu	Gly	Gly	Phe	Tyr	Gly	Gly
				35		40			45						
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Ser	Asp	Leu
				50		55			60						
Gln	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly

65                   70                   75                   80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
              85               90               95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
             100              105              110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
             115              120              125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
             130              135              140  
Leu Thr  
145

<210> 497  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 497  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1               5               10               15  
His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu  
20              25              30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35              40              45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50              55              60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65              70              75              80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85              90              95  
Ile Leu Arg Lys Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100             105             110  
Ser Ala Ser Gly Tyr Tyr Arg Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115             120             125  
Val Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130             135             140  
Leu Thr  
145

<210> 498  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 498  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1               5               10               15  
His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu  
20              25              30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35              40              45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50              55              60

Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 499  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 499  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 500  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 500  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu

50               55               60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65               70               75               80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Gln  
85               90               95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100              105              110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115              120              125  
Val Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130              135              140  
Leu Thr  
145

<210> 501  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
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<400> 501  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1               5               10               15  
His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu  
20              25              30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Gly Gly  
35              40              45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50              55              60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65              70              75              80  
Tyr Arg Asp Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Glu  
85              90              95  
Ile Leu Arg Lys Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100             105             110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115             120             125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Lys  
130             135             140  
Ile Thr  
145

<210> 502  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 502  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
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His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu  
20              25              30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Gly Gly  
35              40              45

Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Asp Gln Lys Ala Gly Ser Thr Leu Ile Lys His Ala Glu Gln  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 503  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 503  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Ile Thr  
145

<210> 504  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 504  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Arg Val Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly

35                   40                   45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50                   55                   60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65                   70                   75                   80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85                   90                   95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100                 105                 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115                 120                 125  
Val Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130                 135                 140  
Leu Thr  
145

<210> 505  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 505  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1                 5                 10                 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Ile Glu Val Cys Met Tyr Glu  
20                 25                 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35                 40                 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50                 55                 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65                 70                 75                 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Glu  
85                 90                 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100                105                110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115                120                125  
Ile Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130                135                140  
Ile Thr  
145

<210> 506  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 506  
Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1                 5                 10                 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Ile Glu Val Cys Met Tyr Glu  
20                 25                 30

Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Phe Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Lys His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Met Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Ile Phe Glu Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 507

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 507

Met Leu Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Leu Arg  
1 5 10 15  
His Lys Ile Leu Arg Pro Asn Gln Pro Leu Glu Val Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Arg Gly Ala Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Asp Leu  
50 55 60  
Gln Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Ser Leu Ile Arg His Ala Glu Glu  
85 90 95  
Ile Leu Arg Lys Arg Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Ala Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Phe Asp Thr Pro Pro Val Gly Pro His Ile Leu Met Tyr Lys Arg  
130 135 140  
Leu Thr  
145

<210> 508

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 508

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Lys Tyr Glu

	20	25	30												
Thr	Asp	Leu	Leu	Gly	Gly	Thr	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
			35		40							45			
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Asn	Ala	Glu	His	Ser	Glu	Leu
			50		55						60				
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
			65		70				75			80			
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
			85				90					95			
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr	
								100	105			110			
Ser	Val	Ser	Gly	Tyr	Tyr	Lys	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
								115	120			125			
Val	Tyr	Asp	Ile	Pro	Pro	Ile	Gly	Pro	His	Ile	Leu	Met	Tyr	Lys	Lys
								130	135			140			
Leu	Thr														
	145														

<210> 509

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 509

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
1				5				10				15			
His	Arg	Ile	Leu	Arg	Pro	Asn	Gln	Pro	Leu	Glu	Ala	Cys	Met	Tyr	Glu
								20	25			30			
Thr	Asp	Leu	Leu	Gly	Gly	Thr	Phe	His	Leu	Gly	Gly	Tyr	Tyr	Arg	Gly
								35	40			45			
Lys	Leu	Ile	Ser	Ile	Ala	Ser	Phe	His	Gln	Ala	Glu	His	Ser	Glu	Leu
								50	55			60			
Glu	Gly	Gln	Lys	Gln	Tyr	Gln	Leu	Arg	Gly	Met	Ala	Thr	Leu	Glu	Gly
								65	70			75		80	
Tyr	Arg	Glu	Gln	Lys	Ala	Gly	Ser	Thr	Leu	Ile	Arg	His	Ala	Glu	Glu
								85	90			95			
Leu	Leu	Arg	Lys	Gly	Ala	Asp	Leu	Leu	Trp	Cys	Asn	Ala	Arg	Thr	
								100	105			110			
Phe	Val	Ser	Gly	Tyr	Tyr	Glu	Lys	Leu	Gly	Phe	Ser	Glu	Gln	Gly	Glu
								115	120			125			
Val	Tyr	Asp	Ile	Pro	Pro	Ile	Gly	Pro	Tyr	Ile	Leu	Met	Tyr	Glu	Lys
								130	135			140			
Leu	Thr														
	145														

<210> 510

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 510

Met	Ile	Glu	Val	Lys	Pro	Ile	Asn	Ala	Glu	Asp	Thr	Tyr	Glu	Ile	Arg
1				5				10				15			

His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Cys Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Ala  
145

<210> 511  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 511  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Pro Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Glu Lys Leu Gly Phe Ser Glu Gln Gly Glu  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 512  
<211> 146  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic Protein Sequence

<400> 512  
Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg

1	5	10	15
His Arg Ile Leu Arg Pro Asn Gln Pro	Leu Glu Ala Cys Lys Tyr Glu		
20	25	30	
Thr Asp Leu Leu Gly Gly Thr Phe His	Leu Gly Gly Tyr Tyr Arg Gly		
35	40	45	
Lys Leu Ile Ser Ile Ala Ser Phe His	Asn Ala Glu His Ser Glu Leu		
50	55	60	
Glu Gly Gln Lys Gln Tyr Gln Leu Arg	Gly Met Ala Thr Leu Glu Gly		
65	70	75	80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr	Leu Ile Arg His Ala Glu Glu		
85	90	95	
Leu Leu Arg Lys Lys Gly Ala Asp	Leu Leu Trp Cys Asn Ala Arg Thr		
100	105	110	
Ser Val Ser Gly Tyr Tyr Lys Lys	Leu Gly Phe Ser Glu Gln Gly Glu		
115	120	125	
Val Tyr Asp Ile Pro Pro Ile Gly Pro	His Ile Leu Met Tyr Lys Lys		
130	135	140	
Leu Thr			
145			

<210> 513

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 513

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg			
1	5	10	15
His Arg Ile Leu Arg Pro Asn Gln Pro	Leu Glu Ala Cys Met Tyr Glu		
20	25	30	
Thr Asp Leu Leu Gly Gly Thr Phe His	Leu Gly Gly Tyr Tyr Arg Gly		
35	40	45	
Lys Leu Ile Ser Ile Ala Ser Phe His	Gln Ala Glu His Pro Glu Leu		
50	55	60	
Glu Gly Gln Lys Gln Tyr Gln Leu Arg	Gly Met Ala Thr Leu Glu Gly		
65	70	75	80
Tyr Arg Glu Gln Lys Ala Gly Ser Thr	Leu Ile Arg His Ala Glu Glu		
85	90	95	
Leu Leu Arg Lys Lys Gly Ala Asp	Leu Leu Trp Cys Asn Ala Arg Thr		
100	105	110	
Ser Val Ser Gly Tyr Tyr Glu Lys	Leu Gly Phe Ser Glu Gln Gly Glu		
115	120	125	
Val Cys Asp Ile Pro Pro Ile Gly Pro	His Ile Leu Met Tyr Lys Lys		
130	135	140	
Leu Thr			
145			

<210> 514

<211> 146

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Protein Sequence

<400> 514

Met Ile Glu Val Lys Pro Ile Asn Ala Glu Asp Thr Tyr Glu Ile Arg  
1 5 10 15  
His Arg Ile Leu Arg Pro Asn Gln Pro Leu Glu Ala Cys Met Tyr Glu  
20 25 30  
Thr Asp Leu Leu Gly Gly Thr Phe His Leu Gly Gly Tyr Tyr Arg Gly  
35 40 45  
Lys Leu Ile Ser Ile Ala Ser Phe His Gln Ala Glu His Ser Glu Leu  
50 55 60  
Glu Gly Gln Lys Gln Tyr Gln Leu Arg Gly Met Ala Thr Leu Glu Gly  
65 70 75 80  
Tyr Arg Glu Gln Lys Ala Gly Ser Thr Leu Ile Arg His Ala Glu Glu  
85 90 95  
Leu Leu Arg Lys Gly Ala Asp Leu Leu Trp Cys Asn Ala Arg Thr  
100 105 110  
Ser Val Ser Gly Tyr Tyr Lys Lys Leu Gly Phe Ser Glu Gln Gly Gly  
115 120 125  
Val Tyr Asp Ile Pro Pro Ile Gly Pro His Ile Leu Met Tyr Lys Lys  
130 135 140  
Leu Thr  
145

<210> 515

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA Sequence

<400> 515

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